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Lacourciere, Karen

Sent:

Thursday, January 09, 2003 12:38 PM STIC-Biotech/ChemLib

To:

Subject:

sequence search request 09/757,100

Please search SEQ ID NO:18 of 09/757,100 in both the commercial databases and the pending files (Interference)

Thank-you!-

Karen A. Lacourciere Ph.D.

CM1 11D09 GAU 1635

(703) 208-7523 mailbox 11E12

Point of Contact: Beverly Shears Technical Info. Specialist CM1 1E05 Tel: 308-4994

Searcher:_ Phone: Location: Date Picked Up: Date Completed:_ Searcher Prep/Review:__ Clerical:_ Online time:

TYPE OF SEARCH:
NA Sequences:
AA Sequences:
Structures:
Bibliographic:
Litigation:
Full text:
Patent Family:
Other:

ENDOR/COST (when	e applic.)
STN:	
DIALOG:	
Questel/Orbit:	
DRLink:	
Lexis/Nexis:	
Sequence Sys.:	
WWW/Internet:	
Other (specify):	

U.S. DEPARTMENT OF COMMERCE Patent and Trademark Office

SEARCH REQUEST FORM

Requestor's Name:		Serial Number:							
Date:	Phone:	Art Unit:							
Search Topic: Please write a detailed statement of search that may have a special meaning. Give exar a copy of the sequence. You may include	mples or relevant citations, author	s keywords, etc., if known. For sea	ned. Define any terms quences, please attach						
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PTO-1590 (9-90)

USCOMM-DC 90-3952

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(FILE 'REGISTRY' ENTERED AT 10:55:53 ON 13 JAN 2003) L1 20 S TTTCAACCAGATGGTCATTC/SQSN

FILE 'HCAPLUS' ENTERED AT 10:57:04 ON 13 JAN 2003 L2 7 S L1

ANSWER 1 OF 7 HCAPLUS COPYRIGHT 2003 ACS 1.2 2002:889441 HCAPLUS

ACCESSION NUMBER: DOCUMENT NUMBER:

INVENTOR(S):

137:381006

TITLE:

Expressed sequence profiles and their use for the therapy and diagnosis of human lung cancer Henderson, Robert A.; Wang, Tongtong; Watanabe,

Yoshihiro; Johnson, Jeffrey C.; Retter, Marc W.; Durham, Margarita; Carter, Darrick; Fanger, Gary R.; Vedvick, Thomas S.; Bangur, Chaitanya S.;

McNabb, Andria

PATENT ASSIGNEE(S):

Corixa Corporation, USA

SOURCE:

U.S. Pat. Appl. Publ., 82 pp., Cont.-in-part of

U.S. Ser. No. 849,626.

CODEN: USXXCO

DOCUMENT TYPE:

Patent

LANGUAGE:

English

FAMILY ACC. NUM. COUNT:

PATENT INFORMATION:

	PATENT NO.	KIND	DATE		APPLICATION NO).	DATE
PRIO	US 2002172952 US 6504010 US 2002168637 US 2002197669 RITY APPLN. INFO.	A1 B1 A1 A1:	20021121 20030107 20021114 20021226	US US US US US US US US US US US US US U	1999-346492 1999-466867 1999-476300 2000-519642 2000-533077 2000-560406 2000-589184 2000-614124 2000-651563 2000-658824 2000-671325 2000-677419 2000-702705 2000-736457 2001-849626	A2 A2 A2 A2 A2 A2 A2 A2 A2 A2 A2 A2 A2 A	20010710 20001030 20001213 20010503 19990630 19991015 19991217 19991230 20000306 20000322 20000410 20000427 20000605 20000711 20000829 20000908 20000926 20001006 20001030 20001213 20010503 20000630
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AB Nucleic acid and encoded protein compns. and methods for the therapy and diagnosis of cancer, particularly lung cancer, are disclosed. Thus, cDNAs encoding tumor-assocd. proteins were identified by conventional and PCR-based library subtraction, cDNA microarray anal., and real-time PCR using cDNA libraries isolated from lung adenocarcinoma, small cell lung carcinoma, squamous cell lung carcinoma, and lung neuroendocrine carcinoma vs. a pool of poly(A)+ RNA from 9 normal tissues. Illustrative compns. comprise one or more lung tumor polypeptides, immunogenic portions thereof,

> Searcher : 308-4994 Shears

polynucleotides that encode such polypeptides, antigen presenting cell that expresses such polypeptides, and T cells that are specific for cells expressing such polypeptides. The disclosed compns. are useful, for example, in the diagnosis, prevention and/or treatment of diseases, particularly lung cancer.

IT 475864-34-7

RL: ANT (Analyte); BSU (Biological study, unclassified); DGN (Diagnostic use); PRP (Properties); THU (Therapeutic use); ANST (Analytical study); BIOL (Biological study); USES (Uses) (nucleotide sequence; expressed sequence profiles and their use for the therapy and diagnosis of human lung cancer)

L2ANSWER 2 OF 7 HCAPLUS COPYRIGHT 2003 ACS ACCESSION NUMBER: 2002:869469 HCAPLUS

DOCUMENT NUMBER: 137:364446

TITLE: Expressed sequence profiles and their use for the therapy and diagnosis of human lung cancer

INVENTOR(S): Wang, Tongtong; Bangur, Chaitanya S.; Lodes,

Michael J.; Fanger, Gary R.; Vedvick, Thomas S.; Carter, Darrick; Retter, Marc W.; Mannion, Jane;

Fan, Liqun; Wang, Aijun

USA PATENT ASSIGNEE(S):

SOURCE: U.S. Pat. Appl. Publ., 66 pp., Cont.-in-part of

U. S. Ser. No. 702,705.

CODEN: USXXCO

DOCUMENT TYPE:

Patent English LANGUAGE:

FAMILY ACC. NUM. COUNT:

PATENT INFORMATION:

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                                                          DATE
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US 2002168637
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        NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR,
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             MD, RU, TJ, TM
         RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH,
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             TG
PRIORITY APPLN. INFO .:
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                                                          A2 19990630
                                        US 1999-419356
                                                          A2 19991015
                                        US 1999-466867
                                                          A2 19991217
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                                                          A2 19991230
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                                                         A2 20000630
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                                        US 2000-658824
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                                        US 2000-702705
                                                          A2 20001030
                                                          A2 20001213
                                        US 2000-736457
                                        US 2001-849626
                                                          A2 20010503
AB
     Nucleic acid and encoded protein compns. and methods for the therapy
     and diagnosis of cancer, particularly lung cancer, are disclosed.
     Thus, cDNAs encoding tumor-assocd. proteins were identified by
     conventional and PCR-based library subtraction, cDNA microarray
     anal., and real-time PCR using cDNA libraries isolated from lung
     adenocarcinoma, small cell lung carcinoma, squamous cell lung
     carcinoma, and lung neuroendocrine carcinoma vs. a pool of poly(A)+
     RNA from 9 normal tissues. Illustrative compns. comprise one or
     more lung tumor polypeptides, immunogenic portions thereof,
     polynucleotides that encode such polypeptides, antigen presenting
     cell that expresses such polypeptides, and T cells that are specific
     for cells expressing such polypeptides. The disclosed compns. are
     useful, for example, in the diagnosis, prevention and/or treatment
     of diseases, particularly lung cancer.
IT
     475019-40-0
     RL: ANT (Analyte); BSU (Biological study, unclassified); DGN
     (Diagnostic use); PRP (Properties); THU (Therapeutic use); ANST
     (Analytical study); BIOL (Biological study); USES (Uses)
        (nucleotide sequence; expressed sequence profiles and their use
        for the therapy and diagnosis of human lung cancer)
L2
     ANSWER 3 OF 7 HCAPLUS COPYRIGHT 2003 ACS
ACCESSION NUMBER:
                         2002:332807 HCAPLUS
DOCUMENT NUMBER:
                         136:351185
TITLE:
                         The contribution of 700,000 ORF sequence tags to
                         the definition of the human transcriptome
AUTHOR(S):
                         Camargo, Anamaria A.; Samaia, Helena P. B.;
                         Dias-Neto, Emmanuel; Simao, Daniel F.; Migotto,
                         Italo A.; Briones, Marcelo R. S.; Costa,
                         Fernando F.; Nagai, Maria Aparecida;
                         Verjovski-Almeida, Sergio; Zago, Marco A.;
                         Andrade, Luis Eduardo C.; Carrer, Helaine;
                         El-Dorry, Hamza F. A.; Espreafico, Enilza M.;
```

Searcher: Shears 308-4994

Habr-Gama, Angelita; Giannella-Neto, Daniel;

Goldman, Gustavo H.; Gruber, Arthur; Hackel, Christine; Kimura, Edna T.; Maciel, Rui M. B.; Marie, Suely K. N.; Martins, Elizabeth A. L.; Nobrega, Marina P.; Paco-Larson, Maria Luisa; Pardini, Maria Ines M. C.; Pereira, Goncalo G.; Pesquero, Joao Bosco; Rodrigues, Vanderlei; Rogatto, Silvia R.; Da Silva, Ismael D. C. G.; Sogayar, Mari C.; Sonati, Maria De Fatima; Tajara, Eloiza H.; Valentini, Sandro R.; Alberto, Fernando L.; Amaral, Maria Elisabete J.; Aneas, Ivy; Arnaldi, Liliane A. T.; De Assis, Angela M.; Bengtson, Mario Henrique; Bergamo, Nadia Aparecida; Bombonato, Vanessa; De Camargo, Maria E. R.; Canevari, Renata A.; Carraro, Dirce M.; Cerutti, Janete M.; Correa, Maria Lucia C.; Correa, Rosana F. R.; Costa, Maria Cristina R.; Curcio, Cyntia; Hokama, Paula O. M.; Ferreira, Ari J. S.; Furuzawa, Gilberto K.; Gushiken, Tsieko; Ho, Paulo L.; Kimura, Elza; Krieger, Jose E.; Leite, Luciana C. C.; Majumder, Paromita; Marins, Mozart; Marques, Everaldo R.; Melo, Analy S. A.; Melo, Monica; Mestriner, Carlos Alberto; Miracca, Elisabete C.; Miranda, Daniela C.; Nascimento, Ana Lucia T. O.; Nobrega, Francisco G.; Ojopi, Elida P. B.; Pandolfi, Jose Rodrigo C.; Pessoa, Luciana G.; Prevedel, Aline C.; Rahal, Paula; Rainho, Claudia A.; Reis, Eduardo M. R.; Ribeiro, Marcelo L.; Da Ros, Nancy; De Sa, Renata G.; Sales, Magaly M.; Sant'anna, Simone Cristina; Dos Santos, Mariana L.; Da Silva, Aline M.; Da Silva, Neusa P.; Silva, Wilson A., Jr.; Da Silveira, Rosana A.; Sousa, Josane F.; Stecconi, Daniella; Tsukumo, Fernando; Valente, Valeria; Soares, Fernando; Moreira, Eloisa S.; Nunes, Diana N.; Correa, Ricardo G.; Zalcberg, Heloisa; Carvalho, Alex F.; Reis, Luis F. L.; Brentani, Ricardo R.; Simpson, Andrew J. G.; De Souza, Sandro J.

CORPORATE SOURCE:

SOURCE:

Ludwig Institute for Cancer Research, Sao Paulo, 01509-010, Brazil

of the United States of America (2001), 98(21),

Proceedings of the National Academy of Sciences

12103-12108 CODEN: DNASA6: ISSN: 0027-8424

CODEN: PNASA6; ISSN: 0027-8424 National Academy of Sciences

PUBLISHER:
DOCUMENT TYPE:
LANGUAGE:

Journal English

AB Open reading frame expressed sequences tags (ORESTES) differ from conventional ESTs by providing sequence data from the central protein coding portion of transcripts. A total of 696,745 ORESTES sequences were generated from 24 human tissues and a subset of the data that correspond to a set of 15,095 full-length mRNAs used as a means of assessing the efficiency of the strategy and its potential contribution to the definition of the human transcriptome. It was estd. that ORESTES sampled over 80% of all highly and moderately expressed, and between 40% and 50% of rarely expressed, human genes. In the most thoroughly sequenced tissue, the breast, the 130,000

Searcher: Shears 308-4994

ORESTES generated are derived from transcripts from an estd. 70% of all genes expressed in that tissue, with an equally efficient representation of both highly and poorly expressed genes. respect, the capacity of the ORESTES strategy both for gene discovery and shotgun transcript sequence generation significantly exceeds that of conventional ESTs. The distribution of ORESTES is such that many human transcripts are now represented by a scaffold of partial sequences distributed along the length of each gene product. The exptl. joining of the scaffold components, by reverse transcription-PCR, represents a direct route to transcript finishing that may represent a useful alternative to full-length cDNA cloning. [This abstr. record is one of many records for this document necessitated by the large no. of index entries required to fully index the document and publication system constraints.].

270220-29-6, GenBank AW992843 270220-33-2, GenBank IT AW992847

RL: BSU (Biological study, unclassified); PRP (Properties); BIOL (Biological study)

(nucleotide sequence; contribution of 700,000 ORF sequence tags to the definition of the human transcriptome)

L2 ACCESSION NUMBER:

DOCUMENT NUMBER:

TITLE:

AUTHOR(S):

ANSWER 4 OF 7 HCAPLUS COPYRIGHT 2003 ACS 2002:332795 HCAPLUS

136:351174

The contribution of 700,000 ORF sequence tags to the definition of the human transcriptome Camargo, Anamaria A.; Samaia, Helena P. B.; Dias-Neto, Emmanuel; Simao, Daniel F.; Migotto, Italo A.; Briones, Marcelo R. S.; Costa, Fernando F.; Nagai, Maria Aparecida; Verjovski-Almeida, Sergio; Zago, Marco A.; Andrade, Luis Eduardo C.; Carrer, Helaine; El-Dorry, Hamza F. A.; Espreafico, Enilza M.; Habr-Gama, Angelita; Giannella-Neto, Daniel; Goldman, Gustavo H.; Gruber, Arthur; Hackel, Christine; Kimura, Edna T.; Maciel, Rui M. B.; Marie, Suely K. N.; Martins, Elizabeth A. L.; Nobrega, Marina P.; Paco-Larson, Maria Luisa; Pardini, Maria Ines M. C.; Pereira, Goncalo G.; Pesquero, Joao Bosco; Rodrigues, Vanderlei; Rogatto, Silvia R.; Da Silva, Ismael D. C. G.; Sogayar, Mari C.; Sonati, Maria De Fatima; Tajara, Eloiza H.; Valentini, Sandro R.; Alberto, Fernando L.; Amaral, Maria Elisabete J.; Aneas, Ivy; Arnaldi, Liliane A. T.; De Assis, Angela M.; Bengtson, Mario Henrique; Bergamo, Nadia Aparecida; Bombonato, Vanessa; De Camargo, Maria E. R.; Canevari, Renata A.; Carraro, Dirce M.; Cerutti, Janete M.; Correa, Maria Lucia C.; Correa, Rosana F. R.; Costa, Maria Cristina R.; Curcio, Cyntia; Hokama, Paula O. M.; Ferreira, Ari J. S.; Furuzawa, Gilberto K.; Gushiken, Tsieko; Ho, Paulo L.; Kimura, Elza; Krieger, Jose E.; Leite, Luciana C. C.; Majumder, Paromita; Marins, Mozart; Marques, Everaldo R.; Melo, Analy S. A.; Melo, Monica; Mestriner, Carlos Alberto; Miracca, Elisabete C.; Miranda, Daniela C.; Nascimento, Ana Lucia

Searcher : Shears 308-4994

T. O.; Nobrega, Francisco G.; Ojopi, Elida P. B.; Pandolfi, Jose Rodrigo C.; Pessoa, Luciana G.; Prevedel, Aline C.; Rahal, Paula; Rainho, Claudia A.; Reis, Eduardo M. R.; Ribeiro, Marcelo L.; Da Ros, Nancy; De Sa, Renata G.; Sales, Magaly M.; Sant'anna, Simone Cristina; Dos Santos, Mariana L.; Da Silva, Aline M.; Da Silva, Neusa P.; Silva, Wilson A., Jr.; Da Silveira, Rosana A.; Sousa, Josane F.; Stecconi, Daniella; Tsukumo, Fernando; Valente, Valeria; Soares, Fernando; Moreira, Eloisa S.; Nunes, Diana N.; Correa, Ricardo G.; Zalcberg, Heloisa; Carvalho, Alex F.; Reis, Luis F. L.; Brentani, Ricardo R.; Simpson, Andrew J. G.; De Souza, Sandro J.

CORPORATE SOURCE:

Ludwig Institute for Cancer Research, Sao Paulo,

01509-010, Brazil

SOURCE:

Proceedings of the National Academy of Sciences of the United States of America (2001), 98(21),

12103-12108

CODEN: PNASA6; ISSN: 0027-8424 National Academy of Sciences

PUBLISHER: DOCUMENT TYPE:

Journal LANGUAGE: English

Open reading frame expressed sequences tags (ORESTES) differ from AB conventional ESTs by providing sequence data from the central protein coding portion of transcripts. A total of 696,745 ORESTES sequences were generated from 24 human tissues and a subset of the data that correspond to a set of 15,095 full-length mRNAs used as a means of assessing the efficiency of the strategy and its potential contribution to the definition of the human transcriptome. It was estd. that ORESTES sampled over 80% of all highly and moderately expressed, and between 40% and 50% of rarely expressed, human genes. In the most thoroughly sequenced tissue, the breast, the 130,000 ORESTES generated are derived from transcripts from an estd. 70% of all genes expressed in that tissue, with an equally efficient representation of both highly and poorly expressed genes. In this respect, the capacity of the ORESTES strategy both for gene discovery and shotgun transcript sequence generation significantly exceeds that of conventional ESTs. The distribution of ORESTES is such that many human transcripts are now represented by a scaffold of partial sequences distributed along the length of each gene product. The exptl. joining of the scaffold components, by reverse transcription-PCR, represents a direct route to transcript finishing that may represent a useful alternative to full-length cDNA cloning. [This abstr. record is one of many records for this document necessitated by the large no. of index entries required to fully index the document and publication system constraints.].

ΙT **272727-43-2**, GenBank BE092864

RL: BSU (Biological study, unclassified); PRP (Properties); BIOL (Biological study)

(nucleotide sequence; contribution of 700,000 ORF sequence tags to the definition of the human transcriptome)

ANSWER 5 OF 7 HCAPLUS COPYRIGHT 2003 ACS L2 ACCESSION NUMBER: 2002:51525 HCAPLUS

DOCUMENT NUMBER: 136:129944

TITLE: Human nucleic acid and expressed protein

> Shears 308-4994 Searcher :

HING FACE DEMINA (USPTO)

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compositions and methods for the therapy and
                           diagnosis of lung cancer
                           Wang, Tongtong; Watanabe, Yoshihiro; Henderson,
INVENTOR(S):
                           Robert A.; Johnson, Jeffrey C.; Retter, Marc W.;
                           Marnerakis, Margarita; Carter, Darrick; Fanger,
                           Gary Richard; Vedvick, Thomas S.; Bangur,
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                           Fanger, Neil; Switzer, Ann; McNeill, Patricia
                           D.; Clapper, Jonathan D.
                           Corixa Corporation, USA
PATENT ASSIGNEE(S):
SOURCE:
                           PCT Int. Appl., 223 pp.
                           CODEN: PIXXD2
DOCUMENT TYPE:
                           Patent
LANGUAGE:
                           English
FAMILY ACC. NUM. COUNT:
PATENT INFORMATION:
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                                           WO 2000-US18061 A2 20000630
AB
     Compns. and methods for the therapy and diagnosis of cancer,
     particularly lung cancer, are disclosed. Illustrative compns.
     comprise one or more lung tumor polypeptides, immunogenic portions
     thereof, the cDNA polynucleotides that encode such polypeptides,
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antigen-presenting cell that expresses such polypeptides, and T cells that are specific for cells expressing such polypeptides.

Thus, lung tumor cDNAs are isolated and characterized using conventional cDNA library subtraction, PCR-based subtraction, microarray anal., and lung chip 5. The cDNA libraries are isolated from lung adenocarcinoma, small cell lung carcinoma, lung neuroendocrine carcinoma, and squamous cell lung carcinoma. The immunogenic gene products are expressed in Escherichia coli, and epitope mapping is provided using antibodies one of the lung tumor-specific antigens, designated L552S. The disclosed compns. are useful, for example, in the diagnosis, prevention and/or treatment of diseases, particularly lung cancer.

IT 388152-61-2P

RL: ANT (Analyte); BPN (Biosynthetic preparation); DGN (Diagnostic use); PRP (Properties); THU (Therapeutic use); ANST (Analytical study); BIOL (Biological study); PREP (Preparation); USES (Uses) (nucleotide sequence; human nucleic acid and expressed protein compns. and methods for the therapy and diagnosis of lung cancer)

L2 ANSWER 6 OF 7 HCAPLUS COPYRIGHT 2003 ACS ACCESSION NUMBER: 2001:731174 HCAPLUS

DOCUMENT NUMBER: 135:283933

TITLE: Nucleic acid markers and methods for the

measurement of sensitivity or resistance to agents used in identification, assessment, prevention, and therapy of human cancers

INVENTOR(S): Lillie, James; Brown, Jeffrey L.; Bolt, Andrew;

Van Huffel, Christophe

PATENT ASSIGNEE(S): Millennium Predictive Medicine, Inc., USA

SOURCE: PCT Int. Appl., 223 pp.

CODEN: PIXXD2

DOCUMENT TYPE: Patent LANGUAGE: English

FAMILY ACC. NUM. COUNT: 1

PATENT INFORMATION:

PA:	TENT I	NO.		KI	ND	DATE			A.	PPLI	CATI	ON NO	Э.	DATE		
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WO	WO 2001073430 A2			2	2001	1004		WO 2001-US9202					20010322			
WO	2001	0734	30	A.	3	2002	0411									
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PRIORITY	Y APP	LN.	INFO	.:				1	US 2	000-	1921	00P	Р	2000	0324	
								1	US 2	000-	1970	64P	P	2000	0413	

AB The present invention is directed to the identification of markers that can be used to det. whether cancer cells are sensitive or resistant to a therapeutic agent. The present invention is also directed to the identification of therapeutic targets. The invention features 436 "sensitivity markers". These are markers that are expressed in most or all cell lines that are sensitive to

Searcher: Shears 308-4994

treatment with an agent and which are not expressed (or are expressed at a rather low level) in cells that are resistant to treatment with that agent. The invention also features 3132 "resistance markers". These are markers that are expressed in most or all cell lines that are resistant to treatment with an agent and which are not expressed (or are expressed at a rather low level) in cells that are sensitive to treatment with that agent. PCR-based subtractive hybridization of cDNA libraries from normal and cancerous tissues was used to identify the markers. At least some of the examples relate to sensitivity or resistance to taxol.

IT **191206-39-0**

RL: ANT (Analyte); BOC (Biological occurrence); BSU (Biological study, unclassified); PRP (Properties); THU (Therapeutic use); ANST (Analytical study); BIOL (Biological study); OCCU (Occurrence); USES (Uses)

(resistance marker; nucleic acid markers and methods for the measurement of sensitivity or resistance to agents used in identification, assessment, prevention, and therapy of human cancers)

L2 ANSWER 7 OF 7 HCAPLUS COPYRIGHT 2003 ACS ACCESSION NUMBER: 2001:551387 HCAPLUS

DOCUMENT NUMBER: 135:103496

TITLE: Nucleic acid compositions, kits, and methods for

identification, assessment, prevention, and

therapy of human breast cancer

INVENTOR(S): Lillie, James; Palermo, Adam; Wang, Youzhen;

Steinmann, Kathleen; Elias, Josh

PATENT ASSIGNEE(S): Millennium Predictive Medicine, Inc., USA

SOURCE: PCT Int. Appl., 2674 pp.

CODEN: PIXXD2

DOCUMENT TYPE: Patent LANGUAGE: English

PATENT INFORMATION:

	PATENT NO. KIND					ND I	DATE	•		A	PPLI	CATIO	э.	DATE			
						:											
	WO :	20010)4669	97 A	2	:	2001	0628		M(200	00-08	3352.	14	2000	1221	
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AB The invention relates to nucleic acid marker compns., kits and methods for detecting, characterizing, preventing, and treating human breast cancers. A variety of markers are provided, wherein changes in the levels of expression of one or more of the nucleic acid markers is correlated with the presence of breast cancer. The

Searcher: Shears 308-4994

level of expression of numerous potential markers was measured in cells obtained from breast cancer tissue samples obtained form fifteen patients afflicted with breast cancer and from eleven breast cancer cell cultures, based on comparison with expression levels of each marker in corresponding non-cancerous breast tissue and cell cultures. The 15 cancer tissue samples include (i) five invasive lobular carcinomas (ILC), (ii) five invasive ductal carcinomas (IDC), and (iii) five samples of ductal carcinoma in situ (DCIS). As an addnl. evaluation of ability to indicate breast cancer, individual markers that were identified by transcriptional profiling criteria were also tested in six different subtracted library expts. In addn., protein profiling expts. were undertaken to assess whether the proteins assocd. with the expression of individual markers of the invention are secreted. Table 21 lists approx. 43,500 GenBank Accession Nos. from the present invention. [This abstr. record is two of eight records for this document necessitated by the large no. of index entries required to fully index the document and publication system constraints.].

IT 191206-39-0

RL: BSU (Biological study, unclassified); PRP (Properties); BIOL (Biological study)

(nucleotide sequence; nucleic acid compns., kits, and methods for identification, assessment, prevention, and therapy of human breast cancer)

E61 THROUGH E67 ASSIGNED

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L3 ANSWER 1 OF 7 REGISTRY COPYRIGHT 2003 ACS

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OTHER NAMES:

CN 981: PN: US20020172952 SEQID: 981 claimed DNA

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REFERENCE 1: 137:381006

L3 ANSWER 2 OF 7 REGISTRY COPYRIGHT 2003 ACS

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REFERENCE 1: 137:364446

L3 ANSWER 3 OF 7 REGISTRY COPYRIGHT 2003 ACS

RN **388152-61-2** REGISTRY

Searcher: Shears 308-4994

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Searcher: Shears 308-4994

REFERENCE 2: 135:103496

FILE 'HOME' ENTERED AT 10:58:05 ON 13 JAN 2003

Searcher: Shears 308-4994

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SUMMARIES

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TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0476
CURRENT APPLICATION UNMBER: PCT/US00/18999
CURRENT FILING DATE: 2000-07-13
PRIOR APPLICATION NUMBER: 09/377,310
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PRIOR FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
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LENGTH: 20
TYPE: DNA
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APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
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CURRENT APPLICATION NUMBER: US/09/377,310A
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Gaarde, William A.
Nero, Pamela S.
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1513, Ap
5123, Ap
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6883, Ap
6983, Ap
698
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GENERAL INFORMATION:
APPLICANT: Monia, Brett P.
APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
APPLICANT: Nero, Pamela S.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-053
CURRENT APPLICATION NUMBER: US/09/757,100B
CURRENT FILING DATE: 2001-03-15
PRIOR APPLICATION NUMBER: 09/377,310
PRIOR APPLICATION NUMBER: P07/370,310
PRIOR APPLICATION NUMBER: P07/370,310
PRIOR FILING DATE: 1999-08-19
PRIOR FILING DATE: 1999-08-19
PRIOR FILING DATE: 2000-07-13
NUMBER OF SEQ ID NOS: 44
SOFTWARE: Patentin Ver. 2.0
                                 FILE REFERENCE: 20411-754CON1
CURRENT APPLICATION NUMBER: US/09/332,782
CURRENT FILING DATE: 1999-06-14
EARLIER APPLICATION NUMBER: US 09/181,430
EARLIER FILING DATE: 1998-10-28
NUMBER OF SEQ ID NOS: 21027
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 6610
LENGTH: 325
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NUMBER OF SEQ ID NOS: 43
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 18
LENGTH: 20
TYPE: DNA
TYPE: DNA
FEATURE.
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; OTHER INFORMATION: antisense sequence US-09-377-310-18
                                                                                                                                                                                                                                                                                                  US-09-332-782-6610/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: antisense sequence US-09-757-100B-18
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                                                                                                                                                                                                Sequence 6610, Application US/09332782
GENERAL INFORMATION:
APPLICANT: Hyseq, Inc.
TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED
TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Loc
Matches
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LENGTH: 20
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Best Local Similarity
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ORGANISM: Homo sapiens
                     TYPE: DNA
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20; Conserv
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Conservative 0; Mismatches
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APPLICANT: HOULGATE, REMI
APPLICANT: BIRNBAUM, DANIEL
APPLICANT: BIRNBAUM, DANIEL
APPLICANT: BIRNBAUM, DANIEL
APPLICANT: BIRNBAUM, DANIEL
APPLICANT: VIEW, PATRICE
APPLICANT: FERT, VINCENT
TITLE OF INVENTION: GENE EXPRESSION PROFILING OF PRIMARY BREAST CARCINOMAS
TITLE OF INVENTION: USING ARRAYS OF CANDIDATE GENES
FILE REFERENCE: 1546-R-00
CURRENT APPLICATION NUMBER: US/10/007,926A
PRIOR APPLICATION NUMBER: 60/254,090
PRIOR PILING DATE: 2000-12-08
NUMBER OF SEO ID NOS: 468
SOFTWARE: PATENTIN VET: 2.1
      Gaps
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ORGANISM: Homo sapiens
PEATURE:
OTHER INFORMATION: 5' terminal sequence. ptk2 protein tyrosine
OTHER INFORMATION: kinase 2 (PTK2) gene.
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US-09-359-067-23180/c
Sequence 23180, Application US/09359067
GENERAL INPORMATION:
TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED
TITLE OF INVENTION: FROW VARIOUS CDNA LIBRARIES
TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES
CURRENT APPLICATION NUMBER: US/09/359,067
CURRENT FILING DATE: 1990-07-22
EARLIER RPLING DATE: 1998-08-10
NUMBER OF SEQ ID NOS: 49786
SOFTWARE: FRELEE 1998-08-10
NUMBER OF SEQ ID NOS: 49786
SEQ ID NO 23180
LENGTH: 379
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    Indels
    0
    Mismatches
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; GENERAL INFORMATION:
, APPLICANT: BERTUCCI, FRANCOIS
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LOCATION: (275)
OTHER_INFORMATION: a, t, c
                                        TTTCAACCAGATGGTCATTC
                                                                               TTTCAACCAGATGGTCATTC
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Best Local Similarity 100.
Matches 20; Conservative
    Conservative
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OTHER INFORMATION: a, t,
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LOCATION: (364)
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US-10-007-926A-284/c
  20;
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                                      Score 20; DB 17; Length 325;
Pred. No. 9.5;
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APPLICANT: Wi, James
APPLICANT: Yim, Kenneth
TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED
TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES
FILE REFERENCE: 754CIP
CURRENT APPLICATION NUMBER: US/09/515,694
CURRENT FILING DATE: 2000-02-29
EARLIER APPLICATION NUMBER: 09/332,782
EARLIER FILING DATE: 1999-06-14
EARLIER FILING DATE: 1999-10-18
SARLIER FILING DATE: 1999-10-28
NUMBER OF SEQ ID NOS: 21027
SOFTWARE: FRALESEQ for Windows Version 3.0
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Best Local Similarity 100.0%;
Matches 20; Conservative 0
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APPLICANT: Asghari, Vida
APPLICANT: Blun, Linda
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Sahourieh, Hannah
Sidhu, Navjiwan
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Engleman, Carrie
Faulkner, Brandy
Fox, Melvin
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Lomelli, Michelle
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Smythe, Ashleigh
Tkach, Joe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Garcia, Veronica
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Verna, Ron
Wachter, Adam
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Phuong
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Drake, Jim
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Jesse
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ORGANISM: Homo sapiens
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Best Local Similarity
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US-09-332-782-6610
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US-09-824-557-1277/c
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                                                                                 US-09-790-435-4313
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Query Match
Best Local Similarity 100.
Matches 20; Conservative
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                                                                                                                                                                                                                            SEQ ID NO 4313
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PRIOR APPLICATION NUMBER: US 60/193,438
PRIOR FILING DATE: 2000-03-31
NUMBER OF SEQ ID NOS: 9773
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TITLE OF INVENTION: NOVEL NUCLEIC ACID MOLECULES AND USES
TITLE OF INVENTION: THEREFOR
FILE REFERENCE: 1600.2096-001
CURRENT APPLICATION NUMBER: US/09/824,557
CURRENT FILING DATE: 2001-04-02
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TITLE OF INVENTION: NOVEL NUCLEIC ACID MOLECULES AND USES
TITLE OF INVENTION: THEREFOR
FILE REFERENCE: 1600.2060-001
CURRENT APPLICATION NUMBER: US/09/790,435
CURRENT APPLICATION NUMBER: US/09/790,435
CURRENT FILING DATE: 2001-02-22
PRIOR APPLICATION NUMBER: US 60/183,730
PRIOR FILING DATE: 2000-02-22
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NAME/KEY: misc feature

LOCATION: (1)...(421)

OTHER INFORMATION: n = A,T,C or

-09-824-557-1277
                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 7473
SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                        FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(433)
OTHER INFORMATION: n = A,T,C or
                                                                                                                                                                   LENGTH: 433
TYPE: DNA
ORGANISM: Homo sapiens
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TYPE: DNA
ORGANISM: Homo sapiens
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                      100.0%;
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Pred. No. 10;
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           Mismatches
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                                            Length 433;
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           Indels
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         Gaps
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SEQUENCE 4292, Application US/0939397

GENERAL INFORMATION:
APPLICANT: Hyseq, Inc.
TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED
TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES
FILE REFERENCE: 20411-765
CURRENT FALING DATE: 2001-08-24
PRIOR APPLICATION NUMBER: 09/289,768
PRIOR APPLICATION NUMBER: 09/289,768
PRIOR APPLICATION NUMBER: 09/289,768
PRIOR TILING DATE: 1999-04-08
NUMBER OF SEQ ID NOS: 39996
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 4292
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; NAME/KEY: misc feature
; LOCATION: (1). (436)
; OTHER INFORMATION: n = A,T,C or G
US-09-289-768-4292
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US-09-289-768-4292/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Hyseq, Inc.

APPLICANT: Hyseq, Inc.

TITLE OF INVENTION: ROWEL NUCLEIC ACID SEQUENCES OBTAINED

TITLE OF INVENTION: ROW VARIOUS cDNA LIBRARIES

FILE REFERENCE: 20411-765

CURRENT EPPLICATION NUMBER: US/09/289,768

CURRENT FILING DATE: 1999-04-08

NUMBER OF SEQ ID NOS: 39996

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 4292
                                                                    Query Match
Best Local Similarity
Matches 20; Conserv
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                                                                                                                                                        FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(436)
OTHER INFORMATION: n = A,T,C
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TYPE: DNA
ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
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Local Similarity 100.0%;
hes 20; Conservative 0
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 TTTCAACCAGATGGTCATTC 78
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                        TTTCAACCAGATGGTCATTC 20
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Pred. No.
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Pred. No.
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                                                                    Pred. No. 10;
Mismatches
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Mismatches
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                                                                                                        Length 436;
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                                                                    Gaps
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US-09-431-517-3422/c ; Sequence 3422, Application US/09431517

RESULT 12

Gaps

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100.0%; Score 20; DB 34; Length 449; 100.0%; Pred. No. 10; cive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 15
US-09-824-557-3555/c
; Sequence 3555, Application US/09824557
; GENERAL INFORMATION:
; APPLICANT: Shyjan, Andrew W.
; TITLE OF INVENTION: HOVEL NUCLEIC ACID MOLECULES AND USES
; TITLE OF INVENTION: HOWER: US/09/824,557
; CURRENT APPLICATION NUMBER: US/09/824,557
; CURRENT FILING DATE: 2001-04-02
; PRIOR APPLICATION NUMBER: US 60/193,438
; NUMBER OF SEQ ID NOS: 9773
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 3555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 20; DB 31;
100.0%; Pred. No. 10;
tive 0; Mismatches 0;
CURRENT APPLICATION NUMBER: US/09/909,629
CURRENT FILING DATE: 2001-07-19
PRIOR APPLICATION NUMBER: 09/306,350
PRIOR FILING DATE: 1999-05-07
NUMBER OF SEQ ID NOS: 41304
SOFTWARE: Dt. CT_1 Version 1.1
SEQ ID NO 30669
LENGTH: 449
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            completed: January 12, 2003, 12:04:06
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OTHER INFORMATION: n = A,T,C or G
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Matches 20; Conservative
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Matches 20; Conservative
                                                                                                                                                                                         TYPE: DNA
ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
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Job time: 3167 secs
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100.0%; Score 20; DB 18; Length 443;
Best Local Similarity 100.0%; Pred. No. 10;
Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Dickoon, Mark C.

APPLICANT: Dickoon, Mark C.

APPLICANT: Dranac, Radoje T.

APPLICANT: Jones, Lee W.

APPLICANT: Stache-Crain, Birgit

TITLE OF INVENTION: Novel Nucleic Acid Sequences Obtained

TITLE OF INVENTION: From Various CDNA Libraries

FILE REFERENCE: 20411-776

CURRENT APPLICATION DROPE: US/09/306,350A

CURRENT PILLION DATO: 1999-05-07

NUMBER OF SEQ ID NOS: 41304

SEQ ID NO 30669

LENGTH: 449
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 100.0%; Score 20; DB 17; Length 449; Best Local Similarity 100.0%; Pred. No. 10; Matches 20; Conservative 0; Mismatches 0; Indels
            APPLICANT: Hyseq, Inc
TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED
TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES
FILE REFERENCE: 20411-751CON1
CURRENT APPLICATION NUMBER: US/09/431,517
CURRENT FILING DATE: 1999-11-01
EARLIER APPLICATION NUMBER: US 09/170,294
EARLIER FILING DATE: 1998-10-13
NUMBER OF SEQ ID NOS: 31760
SEQ ID NO 3422
LENGTH: 443
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GENERAL INFORMATION:
APPLICANT: Dramanac, Radoje T.
APPLICANT: Jones, Lee W.
APPLICANT: Labat, Ivan
APPLICANT: Stache-Crain, Birgit
TITLE OF INVENTION: Novel Nucleic Acid Sequences Obtained
TITLE OF INVENTION: From Various cDNA Libraries
FILE REFERENCE: 20411-776
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-306-350A-30669/c; Sequence 30669, Application US/09306350A; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: misc_feature

| LOCATION: (1)...(443)

| OTHER INFORMATION: n = A,T,C or G

US-09-431-517-3422
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       73 TTTCAACCAGATGGTCATTC 54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 TTTCAACCAGATGGTCATTC 20
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US-09-306-350A-30669
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US-09-909-629-30669/c
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100.0%; Score 20; DB 1; Length 550; 100.0%; Pred. No. 0.47; ive 0; Mismatches 0; Indels
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Watanabe, Yoshihiro
Kalos, Michael D.
Sleath, Paul R.
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APPLICANT: Wang, Tongtong
APPLICANT: Watanabe, Yoshihiro
APPLICANT: Kalos, Michael D.
APPLICANT: Sleath, Paul R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      rrrcaaccagargercarrc 214
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Best Local Similarity 100.
Matches 20; Conservative
; ORGANISM: Homo sapiens
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US-10-283-017-981
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                      TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20758, A Sequence 20748, A Sequence 20748, A Sequence 92, Appl Sequence 20760, A
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Sequence 981, App
                                                                                                                                             ; Search time 168 Seconds
(without alignments)
110.212 Million cell updates/sec
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1: /cgn2 6/ptodata/1/pna/PCT_NEW_COMB.seg:*

2: /cgn2_6/ptodata/1/pna/USO6_NEW_COMB.seg:*

3: /cgn2_6/ptodata/1/pna/USO9_NEW_COMB.seg:*

4: /cgn2_6/ptodata/1/pna/USO9_NEW_COMB.seg:*

5: /cgn2_6/ptodata/1/pna/USO9_NEW_COMB.seg:*

6: /cgn2_6/ptodata/1/pna/USO9_NEW_COMB.seg:*

7: /cgn2_6/ptodata/1/pna/USO9_NEW_COMB.seg:*
                  GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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US-09-724-676A-20756
US-09-724-676A-20755
US-09-724-676A-20755
US-09-724-676A-20750
US-09-724-676A-20750
US-09-724-676A-20750
US-09-724-676A-20752
US-09-724-676A-20751
US-09-724-676-20751
US-09-724-676-20754
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18-09-724-676-20754

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18-09-724-676-20758
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US-09-724-676-20760
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Maximum Match 100%
Listing first 45 summaries
                                                                                                     OM nucleic - nucleic search, using sw model
                                                                                                                                                  January 12, 2003, 09:51:30
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Gapop 10.0 , Gapext 1.0
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Maximum DB seq length: 200000000
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222, Appl
379126, A
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Sequence
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APPLICANT: Henderson, Robert A.
APPLICANT: Henderson, Robert A.
APPLICANT: Watanabe, Yoshihiro
APPLICANT: Kalos, Michael D.
APPLICANT: Sleath, Paul R.
APPLICANT: Sleath, Paul R.
APPLICANT: Retter, Marc W.
APPLICANT: Panger, Carrer, Darrick
APPLICANT: Fanger, Cary R.
APPLICANT: Bengur, Chaitanya S.
APPLICANT: Carrer, Darrick
APPLICANT: McNabb, Andria
APPLICANT: McNabb, Andria
TITLE OF INVENTION: CMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210.14.7803PC
CURRENT APPLICATION NUMBER: PCT/USO2/34777
CURRENT FILING DATE: 2002-10-28
NUMBER OF SEQ ID NOS: 2157
SOFTWARER FREESE FREE
US-09-724-676A-20760
US-10-203-138A-10274
US-10-203-138A-10274
US-10-196-793A-22
US-09-513-995-18126
US-09-531-113-37959
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US-09-724-676-5046
US-09-724-676-5040
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Sequence 20756, Application US/09724676A
GENERAL INFORMATION:
APPLICANT: Compugen LTD
TITLE OF INVENTION: Variants of alternative spl
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676A
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
SOFTWARE: PatentIn version 3.2
LENGTH: 1317
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; SEQ ID NO 20756
; LENGTH: 1317
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-724-676-20756
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 981
LENGTH: 550
TYPE: DNA
ORGANISM: Homo sapiens
US-10-283-017-981
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-724-676A-20756
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/10/283,017
CURRENT FILING DATE: 2002-10-28
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les 20; Conserv
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Bangur, Chaitanya S.
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Pred. No. 0.47;
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RESULT 7
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US-09-724-676-20750/c
Commence 20750, Application US/09724676
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CURRENT APPLICATION NUMBER: US/09/724,676
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
SOFTWARE: PatentIn version 3.2
SEQ ID NO 20755
                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn version 3.2 SEQ ID NO 20755
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SEQ ID NO 20750
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             APPLICANT: Compugen LTD
TITLE OF INVENTION: Variants of alternative
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676
CURRENT FILLING DATE: 2000-11-28
NUMBER OF SEO ID NOS: 97222
SOFTWARE: PatentIn version 3.2
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TITLE OF INVENTION: Variants of alternative
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676A
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
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TYPE: DNA
ORGANISM: Homo sapiens
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TYPE: DNA
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                                                                                                                                                                                                                  100.0%; Score 20; DB 5; Length 1836; 100.0%; Pred. No. 0.56; ive 0; Mismatches 0; Indels
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US-09-724-676A-20752/c
is Sequence 20752/c
sequence 20752, Application US/09724676A
sequence 20752, Application US/09724676A
sequence 20752, Application US/09724676A
refulcaNT: Compugen LTD
refulcaNT: Compugen LTD
refulcaNT: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676A
cURRENT FILING DATE: 2000-11-28
number OF SEQ ID NOS: 97222
sequence 2000-11-28
sequence 2000-11-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Compugen LTD
TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
SOFTWARE: PatentIn version 3.2
SOFTWARE: A100
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Pred. No. 0.62;
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CURRENT APPLICATION NUMBER: US/09/724,676A CURRENT FILING DATE: 2000-11-28 NUMBER OF SEQ ID NOS: 97222 SOFTWARE: Patentin version 3.2 SEQ ID NO 20749 LENGTH: 1836
                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-724-676-20752/c
; Sequence 20752, Application US/09724676
; GENERAL INFORMATION:
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100.0%; Pr
tive 0;
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Matches 20; Conservative
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Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                    ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-724-676A-20749
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; ORGANISM: Homo sapiens
US-09-724-676A-20752
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US-09-724-676-20752
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US-09-724-676-20751/c
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LENGTH: 3610
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100.0%; Score 20; DB 5; Length 1791;
Best Local Similarity 100.0%; Pred. No. 0.56;
Matches 20; Conservative 0; Mismatches 0; Indels
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Sequence 20750, Application US/09724676A

GENERAL INFORMATION

TITLE OF INVENTION: Variants of alternative splicing

TITLE OF INVENTION: Variants of alternative splicing

FILE REFERENCE: 129181-4 Compugen

CURRENT FILING DATE: 2000-11-28

NUMBER OF SEQ ID NOS: 97222

SOFTWARE: PatentIn version 3.2

SEQ ID NO 20750

LENGTH: 1791
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Sequence 20749, Application US/09724676

GENERAL INFORMATION:

TITLE OF INVENTION:

FILE REFERENCE: 129181.4 Compugen

CURRENT APPLICATION NUMBER: US/09/724,676

CURRENT FILING DATE: 2000-11-28

NUMBER OF SEQ ID NOS: 97222

SEQ ID NO 20749

LENGTH: 1836
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TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129181.4 Compugen
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ilarity 100.0%; Pred. No. 0.56;
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; Sequence 20749, Application US/09724676A
; GENERAL INFORMATION:
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Best Local Similarity 100.0
Matches 20; Conservative
                                          ; ORGANISM: Homo sapiens
US-09-724-676-20750
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US-09-724-676A-20750
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; ORGANISM: Homo sapiens
US-09-724-676-20749
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Best Local Similarity
Matches 20; Conserv
  LENGTH: 1791
                          TYPE: DNA
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Sequence 20751, Application US/09724676
GENERAL INFORMATION:
APPLICANT: Compugen LTD
TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
SOFTWARE: Patentin version 3.2
SEQ ID NO 20751
LENGTH: 3677
TYDE: DNA
ORGANISM: Homo sapiens
US-09-724-676-20751
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US-09-724-676-20754/c
Sequence 20754, Application US/09724676
GENERAL INFORMATION:
APPLICANT: Compugen LTD
TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
SOFTWARE: PatentIn version 3.2
SEQ ID NO 20754
LENGTH: 3697
TYPE: DNA
ORGANISM: Homo sapiens
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US-09-724-676A-20751/c
; Sequence 20751, Application US/09724676A
; GENERAL INFORMATION:
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; ORGANISM: Homo sapiens
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TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676A
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
SOFTWARE: PatentIn version 3.2
SEQ ID NO 20751
LENGTH: 3677
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Search completed: January 12, 2003, 12:07:16 Job time: 170 secs

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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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                                                                                     OM nucleic - nucleic search, using sw model
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Listing first 45 summaries
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Pred. No. is the number of results predicted by chance to have a

score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Description	AR116330 Sequence G06663 human STS W	L05186 Homo sapien	BC028733 Homo sapi AK094999 Homo sapi	AR116314 Sequence	AL832961 Homo sapi	AX281683 Sequence AC067931 Homo sapi	AJ132004 Rhizobium	ALS91/90 SINOTHIZO AC102801 Homo sapi	AC084336 Homo sapi	ACUII632 HOMO SAPI AE005765 Caulobact	AC103694 Homo sapi	AL/13916 Mouse DNA AC091767 Homo sapi	AL606668 Oryza sat	AC108406 Mus muscu	AC011777 Homo sapi	AC106524 Rattus no	AC094333 Rattus no	AP002851 Homo sapi	AC105614 Rattus no	AL663072 Mouse DNA	AC098455 Rattus no	AB070838 Coriolus AC095594 Rattus no	X54715 M.eugenii m	AX481431 Sequence				ผู้ผู้ผู้	70 Sequence	129			linear PAT 16-MAY-2001				kinase expression	
ді в	AR116330 6 AR116330 6 AX3683 6 AX368371	HUMFAK		AR11631					9 AC084336			_								0								10633		AR06942	ALIGNMENTS		20 bp DNA	n			l adhesion	-A 16 1/-OC1-2000; Qualifiers
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Primer B: AGTTTAGGAATTAAGATGTCL
STS size: 350
PCR Profile:
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                                                                                                   Prepared with primer pairs derived Location/Qualifiers
                                                                                                                                                                                                                                                                                      Protocol
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Mammalia; Eutheria;
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Homo sapiens STSs derived from sequences in dbEST and
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Primer: each 5 pM
dNTPs: each 4 nM
  117
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KCl: 50 mM
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Total Vol: 20 ul
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llarity 100.0%;
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PCR Cycles: 35
Thermal Cycler:
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Annealing: 56 degrees C
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                                             /organism="Homo sapiens"
/db_xref="taxon:9606"
/map="951 H_5; 805.4 cR
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Pred. No.
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RESULT 3
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AX368271
                                                                                                                                            Andre, E. and Becker-Andre, M. Expression of an N-terminally adhesion kinase in brain
                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates; 1 (bases 1 to 3052)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Wang,T., Watanabe,Y., Henderson,R.A., Johnson,J.C., Retter,M.W. Marnetakis,M., Carter,D., Fanger,G.R., Vedvick,T.S., Bangur,C.S. Mcnabb,A., Fanger,N., Switzer,A., Mcneill,P.D. and Clapper,J.D. Compositions and methods for the therapy and diagnosis of lung
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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20; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent: WO 0204514-A 981 17-JAN-2002; CORIXA CORPORATION (US)
                                                                                                                                 Biochem. Biophys. Res. Commun.
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                                                                                        Location/Qualifiers
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15. .2654
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113 c 121 g
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EC_number="2.7.1.1"
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complete
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Kodaira, H., Furuya, T., Takahashi, M., Kikkawa, E., Omura, Y., Abe, K.,
Kamihara, K., Kateuta, N., Sato, K., Tanikawa, M., Yamazaki, M.,
Sugiyama, T., Irie, R., Otsuki, T., Sato, H., Wakamatsu, A., Ishii, S.,
Yamamoto, J., Isono, Y., Kawai-Hio, Y., Saito, K., Nishikawa, T.,
Kimura, K., Yamashita, H., Matsuo, K., Nakamura, Y., Sekine, M.,
Kikuchi, H., Murakawa, R., Kanehori, K., Takahashi-Fujii, A.,
Oshima, A., Sugiyama, A., Kawakami, B., Suzuki, Y., Sugano, S.,
Nagahari, K., Masuho, Y., Nagai, K. and Isogai, T.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Homo sapiens hippocampus cDNA to mRNA, clone_lib:BRHIP2
clone:BRHIP2012923.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ch 100.0%; Score 20; DB 9; Length 3199; 1. Similarity 100.0%; Pred. No. 22; 20; Conservative 0; Mismatches 0; Indels
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RNGHIYQPVGKRDPRAAPREKRPREAPGHLGSLASLSSPADSYNBGVKLQPQEISFPP
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IPLLARASTHREE BMAQKLLNSDLGELINKMKLDAQQVVWTSLQQEYKKQMLTRAHLANV
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Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
Tosh Library Preparation: Michael J. Brownstein (NHGRI) & Shiraki
COBNA Library Arrayed by: The II.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Institute for Systems Biology
DNA Sequencing by: Institute for Systems Biology
Contact: amadan@systemsbiology.org
Contact: amadan@systemsbiology.org
Anup Madan, Jessica Fahey, Erin Helton, Mark Ketteman, Anuradha
Madan, Stephanie Rodrigues, Amy Sanchez and Michelle Whiting
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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
1 (bases 1 to 3199)
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Submitted (29-APR-2002) National Institutes of Health, Mammalian
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Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
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Best Local Similarity 100.0
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human.

ORGANISM

KEYWORDS SOURCE

AUTHORS TITLE JOURNAL

REMARK

COMMENT

REFERENCE

RESULT 5 BC028733/c

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LOCUS

ACCESSION

VERSION

BASE COUNT ORIGIN

PRI 15-JUL-2002 highly similar

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, Abe, K.,

/organism="Homo sapiens"

source

FEATURES

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AUTHORS
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Sequence 1 from patent US 6133031.
AR116314
AR116314.1 GI:14096636
HUMFAKX 3791 bp mRNA
Human focal adhesion kinase (FAK) mRNA,
L13616
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Monia, B.P. and Gaarde, W.A.
Antisense inhibition of focal adhesion kinase expression
Patent: US 6133031-A 1 17-OCT-2000;
Location/Qualifiers
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849 c 924 g
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EYKKQMLTAAHALAVDAKNLLDVIDQARLKMLGQTRPH"
a 697 c 741 g 837 t
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RGSIDREDGSLQGPIGNQHIYQPVGKPDPAAPPKKPPRPGAPGHLGSLASLSSPADSY
NEGVKLQPQEISPPPTANLDRSNDKVYENVTGLVKAVIEMSSKIQPAPPEEYVPMVKE
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GVCMWEILMHGYKDFQGYKNNDVIGRIENGERLDMPBNCPPTLYSLMTKCWASVDPSRR
PRFTELKAQLSTILEEEKAQQEERMRWESRRQATVSWDSGGSDEAPPKESRGYPSPB
SSEGFYPSPQHMVQTNHYQVSGYPGSHGITAMAGSIYPGQASLLDQTDSWNHRPQEIA
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DVHQGIYMSPENPALAVAIKTCKNCTSDSVREKFLQEALTMRQFDHPHIVKLIGVITE
NPVWIIMELCTLGELRSFLQVRKYSLDLASLILYAYQLSTALAYLESKRFVHRDIAAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /codon_start=1
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/clone_lib="BRHIP2"
/note="cloning_vector: pME18SFL3"
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/db_xref="taxon:9606"
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Mammalia; Eutheria; P
1 (bases 1 to 3791)
Whitney,G.S., Chan,P.
                                                                                                                                                       20;
Homo sapiens mRNA; cDNA
AL832961
AL832961.1 GI:21733549
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Human T and B lymphocytes express a structurally conserved focal adhesion kinase, pp125FAK

DNA Cell Biol. 12 (9), 823-830 (1993)
                                        HSM804272
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Homo sapiens (tissue library: lambda ft
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/gene="FAK"
a 849 c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /translation="maaayldpnlnhtpnSsTKThLGTGMERSPGAMERVLKVFHYFE
SNSEPTTWASIIRHGDATDVRGIIQKIVDSHKVKHVACYGFRLSHLRSEEVHWLHVDM
                                                                                                                                                                                                                                                        AKNLLDVIDQARLKMLGQTRPH"
                                                                                                                                                                                                                                                                      PLLPASTHREIEMAQKLLNSDL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /product="focal adhesion kinase"
/protein_id="AAA58469.1"
/db_xref="GI:439875"
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/db_xref="taxon:9606"
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tissue_lib="lambda ft11"
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Primates;
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                        linear PRI 10-JUL-2
n clone DKFZp66600110)
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3558 TTTCAACCAGATGGTCATTC 3539
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                                                                               AC067931
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This close (DKEZp66600110) is available at the REPD in Berlin.
Please contact the REPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de Further
information about the clone and the sequencing project is available
at http://mips.gsf.de/proj/CDNA/.
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                                                    Mammalia; Eutheria; Primates; Catarrhin; Hominidae; Homo.
Mammalia; Eutheria; Primates; Catarrhin; Hominidae; Homo.
I (bases 1 to 4431)
Ansorge,W., Wirkner,U., Mewes,H.W., Weil,B. and Wiemann,S.
Direct Submission
Submitted (09-JUL-2002) 1, D-85764 Neuherberg, GERMANY
Clone from S. Wiemann, Molecular Genome Analysis; German Cancer Research Center (DKF2); Email s. wiemann@dkf2-heidelberg.de; sequenced by EMBL (European Molecular Biology Laboratories, Heidelberg/Germany) within the cDNA sequencing consortium of the
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    /tissue_type="stomach"
/clone_lib="666 (synonym: hsto2). Vector pSport1; host
DH108; sites Not1 + Sal1"
/dev stage="adult"
4383. .4388
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974 c 1045 g 1136 t
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Sequence 92 from Patent WO0177389.
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/db xref="taxon:9606"
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AXZ81683.1 GI:16608934
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                                                                                                                                                                                                                                                                                            Submitted (24-FEB-1999) Sanjuan J., Microbiologia del Suelo y Sistemas Simbioticos, Estacion Experimental del Zaidin-CSIC, Profesor Albareda 1, E-18008 Granada, SPAIN
                                                                                                                                                                                                                                                                                                                                                                                                                               1 (bases 1 to 3974)
Sanjuan-Pinilla,J.M., Olivares,J. and Sanjuan,J.
The Rhizobium meliloti leuA gene is essential for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sinorhizobium meliloti.
Sinorhizobium meliloti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2-isopropylmalate synthase; clp protease; clpP gene; leuA gene; leuCine biosynthesis; ORF142; ORF83; proteolytic subunit.
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AJ132004.1 GI:4539548
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                                                                                                                                                                                                                                                                                                                                                     Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished
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This sequence will be replaced by the finished sequence as soon as it is available and the accession number will be preserved.

1 206467: contig of 206467 bp in length.

Location/Qualifiers
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------ Project Information
Center project name: L10082
Center clone name: 642_A_1
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ilarity 100.0%;
Conservative 0
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                /codon_start=1
/trans1_table=11
/product="CLP protease proteolytic
/protein_id="CAB39976.1"
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                                                                                                                                  /gene="clpP"
136. .723
                                                                                                                                                                           /db_xref="taxon:382"
136. .723
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/clone lib="RPCI-11 Human Male BAC"
50155 c 48369 g 50320 t 4 4
translation="MRNDDDQEEKKTELPLGKETEANLFKSRSIFIYGTITQELAQKV"
                                                                                                            gene="clpP"
                                                                                                                                                                                                              strain="GR4"
                                                                                                                                                                                                                                     organism="Sinorhizobium meliloti"/
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Sinorhizobium meliloti 1021 complete
AL591790 AL591688
AL591790.1 GI:L5075270
1 (bases 1 to 323450)
Capela,D., Barloy-Hubler,F., Gouzy,J., Bothe,G., Ampe,F., Batut,J., Boistard,P., Becker,A., Boutry,M., Cadieu,E., Dreano,S., Gloux,S., Godrie,T., Goffeau,A., Kahn,D., Kiss,E., Lelaure,V., Masuy,D., Pohl,T., Portretelle,D., Puehler,A., Purnelle,B., Ramsperger,U., Renard,C., Thebault,P., Vandenbol,M., Weidner,S. and Gallbert,F. Analysis of the chromosome sequence of the legume symbiont
                                                                                                                                                                                                                              Sinorhizobium meliloti.
Sinorhizobium meliloti
                                                                                                                                                                             Bacteria; Proteobacteria; alpha subdivision; Rhizobiaceae group; Rhizobiaceae; Sinorhizobium.
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SVDLRDGNQALVDPMGHDRKARMPHLLLDMGFKEIEIGFPSASQTDYDPARWCYEEGN
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ATDAAKMITDMAAKAGGGYRFEYSPASEFTGTELEVALEICNAVIEIVRPTADWKLIIN
LPSTVEMATPNIYADQIEWMCRNLDNRENLIVSLHPHNDRGTGIAATDLGLMAGADRV
EGTLFCNGERTGNVDVVTLALMMFTQGVDFKLDCSDIERIKEYSEYSNQMVIPERHPY
VGELVYTAFSGSHQDAINKGMKAIKQAANKPTWEDYPYLPIDERCKEIHQRFIESYVEQPG
GKGGIAYILQEDYGINLPRNLQIEFREEVQRITDEEGKEIHGRFIESYVEQPG
ARIKFVDHHTYPAGEHKGLRVVAAEITDGGGETRQIEGKGTGPIDGFINALSIYLGIEL
SVADYSEHSLQHGSNAAAIAYVEVEYEYGGGKLFGVGINTNIVAASLEAIVSAAANRVLDV
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YVAAPKEQRLCLPNTRFLLHQPSGGTRGMASDIEIQAREIIKMNERLNRIFSEATGQP
VDKIAKDTDRDYWLGAEEAKAYGLVSRIVTSIAEI"
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aaltrlrIIATKGKADRSDLIGEARELVDSFIQALSASEVTIDSHTGVRALDAMARYG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (codon start=1)
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/transl_table=11
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1194. .2903
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PLREKLAALRQQARDRLGPPVHGIDMKKLMDELWEEGE"
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gene="leuA"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 18.4; DB 1;
Pred. No. 1.5e+02;
); Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       743
                                                                                                                                                                                                                                                                                                                                                                                 DNA
                                                                                                                                                                                                                                                                                                                                                        chromosome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 3974;
                                                                                                                                                                                                                                                                                                                                                                               linear
                                                                                                                                                                                                                                                                                                                                                     segment
                                                                                                                                                                                                                                                                                                                                                                                 BCT 05-JUL-2002
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complement (3024...3740)
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SASHVKKVIFHPKFLPGVTICDPELTVGMPKVITAGTGMDAFAHCLEAYSSPFYHPMS
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gene name confidence: hypothetical
predicted by Goodn usage
predicted by Homology
predicted by FrameD"
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Gene name confidence : hypothetical
predicted by Codon usage
predicted by Homology
predicted by FrameD"
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Gene name confidence : hypothetical
predicted by Codon_usage
predicted by Homology
predicted by FrameD"
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complement (3949, .5094)
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complement (5091. .6476)
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                                                                                                                                                                                                                                                                                               Laboratoire de Biologie Moleculaire des Relations
Plantes-Microorganismes, UMR215-CNRS-INRA, BP27, F-31326 Castanet,
France, Laboratoire de Genetique et Devaloppement UMR6051-CNRS,
Facult de Medecine, 2 avenue du Pr. Leon Bernard, F-35043 Rennes,
France, GATC GmbH, Fritz-Arnold-str. 23, D-78467 Konstanz, Germany,
Universitet Bielefeld, Biologie IV (Genetik) Universiteatstr 25,
Universite Catholique de Louvain, Place Croix du Sud 2, Bte 20,
B-1348 Louvain-la-Neuve, Belgium, Unite de Microbiologie, Faculte
des Sciences Agronomiques de Gembloux, Avenue Marechal Juin 6,
B-5030 Gembloux, Balgium E-mail:Gerome Gouzy@toulouse.inra-fr
http://sequence.toulouse.inra-fr/meliloti.html.
                                                                                                                                                                             Direct Submission
Submitted (26-JUL-2001) Gouzy J., Submitted on behalf of the MELILO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /evidence=not_experimental
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Gene name confidence : hypothetical
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Proc. Natl. Acad. Sci. U.S.A. 98 (17), 9877-9882 (2001)
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predicted by FrameD"
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                                                                                                                           (bases 1 to 323450)
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                                  Proc. Nat
21396507
                                                                                              11481430
                                                                                                                                                       Gouzy, J
                                                                                                                                                    AUTHORS
TITLE
JOURNAL
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                                                                                                                    REFERENCE
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REFERENCE
AUTHORS
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TITLE
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VERSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LOCUS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DEFINITION
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Best Local !
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Birren, B., Linton, L., Nusbaum, C., Lander, E., Ali, A., Allen, N., Anderson, S., Barna, N., Bastien, V., Boguslavkiy, L., Boukhgalter, B., Brown, A., Camarata, J., Campopiano, A., Chang, J., Chazaro, B., Choepel, Y., Colangelo, M., Collins, S., Collymore, A., Cook, A., Cooke, P., DeArellano, K., Dewar, K., Diaz, J.S., Dodge, S., Faro, S., Ferreira, P., FitzHugh, W., Gage, D., Galagan, J., Gardyna, S., Ginde, S., Gord, S., Goyette, M., Graham, L., Grand-Pierre, N., Hagos, B., Heaford, A., Horton, L., Hulme, W., Iliev, I., Johnson, R., Jones, C., Kamat, A., Karatas, A., Kells, C., LaRocque, K.,
                                                                                                                                                                                                                                                                                    Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 119671)
Birren,B., Linton,L., Nusbaum,C. and Lander,E.
Homo sapiens chromosome 8, clone RP11-960P16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AC102801
AC102801.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens chromosome 8, clone RP11-960P16, complete sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AC102801
                                                                                                                                                                                                                                                              Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
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predicted by Homology
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ILVGERMVRAFIEAGVPADVFQNLFLDHDTTAALIAAKSFDFINFTGSVEGGRSIERA
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DAFVEKSVARVSNYKLGHDLDFETTLGPMANKEFAATVRNQVADAISKGARALIDFKL
FPQDDGGAYLAPQVLVDVDHSMEFMREETFGPAVGIMKVKSDAEAIELMNDSRYGLTA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          amino acids, amines, peptides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                KEVREIPHTLREAGEALSGSKMLRAAFGEEVVDHYVHAAEWEQQEYDRRVTDWEVARG
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/transl_table=11
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Pred. No. 1.3e+02;
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RES Birren, B., Linton, L., Nusbaum, C., Lander, E., Ali, A., Allen, N., Anderson, S., Barna, N., Bastien, V., Bloom, T., Boguslavkiy, L., Boukhgalter, B., Brown, A., Comapata, J., Collins, S., Collymore, A., Cook, A., Cook, P., DeArellano, K., Dewar, K., Diaz, J.S., Dodge, S., Faro, S., Ferreira, P., FitzHugh, W., Gage, D., Galagan, J., Gardyna, S., Goid, S., Goyette, M., Graham, L., Grand-Pierre, N., Hagos, B., Horton, L., Hulne, W., Iliev, I., Johnson, R., Jones, C., Kamat, A., Karatas, A., Kells, C., LaRocque, K., Lamazares, R., Landers, T., Lehoczky, J., Levine, R., Lindblad-Toh, K., Liu, G., McCarthy, M., McZewan, P., McKernan, K., Meldrim, J., Marthews, C., McCarthy, M., McZewan, P., McKernan, K., Meldrim, J., Meneus, L., Mihova, T., Mlenga, V., Murphy, T., Naylor, J., Nuylor, C., Nicol, R., Oliver, J., Peterson, K., Phunkhang, P., Pierre, N., Pollara, V., Raymond, C., Retta, R., Rieback, M., Riley, R., Rise, C., Rogov, P., Roman, J., Rosetti, M., Roy, A., Santos, R., Schauer, S., Schupback, R., Senaman, S., Severy, P., Spencer, B., Stange-Thomann, N., Stojanovic, N., Viel, R., Vo, A., Wilson, B., Wu, X., Wyman, D., Ve, W.J., Young, G., Zainoun, J., Zembek, L., Zimmer, A., and Zody, M. Ve, W.J., Young, G., Direct Submission
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MacLean,C., Macdonald,P., Major,J., Marquis,N., Matthews,C.,
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Viel,R., Vo,A., Wilson,B., Wu,X., Wyman,D., Ye,W.J., Young,G.,
Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SP6 end overlaps project AC011632 (L2781) by 63.5 kilobases; overlaps project AC100782 (L19699) by 47.5 kilobases. We are submitting only the middle 119.7 kilobases of the project.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Research, 320 Charles Street, Cambridge, MA 02141, USA
on May 1, 2002 this sequence version replaced gi:17060911.
All repeats were identified using RepeatMasker:
Smit, A.F.A. & Green, P. (1996-1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (01-MAY-2002) Whitehead Institute/MIT Center for Genome
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295. .366
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                                                                                                                                                                                                                                                                                                                        organism="Homo sapiens"
|db_xref="taxon:9606"
                                                                                                                                                                   clone_lib="RPCI-11
                                                                                                                                                                                                                                                                                          chromosome="8"
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repeat_region

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Birren, B., Linton, L., Nusbaum, C., Lander, E., Abraham, H., Allen, N., Anderson, S., Barna, N., Bastien, V., Beda, F., Boguslavkiy, L., Boukhgalter, B., Erom, A., Burkett, G., Campopiano, A., Castle, A., Choepel, Y., Colangelo, M., Collins, S., Collymore, A., Cooke, P., DeArellano, K., Dewar, K., Diaz, J.S., Dodge, S., Ferreira, P., Fitzhugh, W., Gasge, D., Galagan, J., Gardyra, S., Ginde, S., Goyette, M., Graham, L., Grand Pierre, N., Hagos, B., Haford, A., Levine, R., Landers, T., Lehoczky, J., Levine, R., Lieu, C., Liu, G., Macdonald, P., Marquis, N., McCarthy, M., McEwan, P., McKernan, K., Morrow, J., Marquis, N., McCarthy, M., McEwan, C., Peterson, K., Morrow, J., Wurphy, T., Naylor, J., Norman, C.H., O'Connor, T., Morrow, J., Wurphy, T., Naylor, J., Norman, C.H., O'Connor, T., Pisani, C., Pollara, V., Raymond, C., Rieback, M., Riley, R., Sougnez, C., Spencer, B., Stange-Thomann, N., Stojanovic, N.,
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Homo sapiens chromosome 8, clone RP11-625F15, complete sequence.
AC084336
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1 (bases 1 to 156235)

Birren, B., Linton, L., Nusbaum, C. and Lander, E.

Homo sapiens chromosome 8, clone RP11-625F15
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Smit, A.F.A. & Green, P. (1996-1997) http://ftp.genome.washington.edu/RM/RepeatMasker.html	Brown, A., Camarata, J., Campopiano, A., Chang, J., Chazaro, B., Choepel, Y., Colangelo, M., Collins, S., Collyne, A., Cooke, P., DeArellano, K., Dewar, K., Diaz, J.S., Dodge, S., Faro, S., Ginde, S., Gord, S., Goyette, M., Galagan, J., Gardyna, S., Ginde, S., Gord, S., Goyette, M., Galagan, J., Garand-Pierre, N., Hagos, B., Heaford, A., Horton, L., Hulme, W., Iliev, I., Johnson, R., Jones, C., Kamat, A., Karatas, A., Kells, C., LaRocque, K., Lamazares, R., Landers, T., Leboczky, J., Levine, R., Liu, G., MacGarthy, M., McEwan, P., McKernan, K., McPheeters, R., Meldrim, J., McCarthy, M., McEwan, P., McKernan, K., McPheeters, R., Meldrim, J., Weneus, L., Mihova, T., Mlenga, V., Murphy, T., Naylor, J., Nguyen, C., Norman, C.H., O'Connor, T., O'Donnell, P., O'Neil, D., Oliver, J., Peterson, K., Phunkhang, P., Pierre, N., Pollara, V., Raymond, C., Retta, R., Rieback, M., Riley, R., Rise, C., Rogov, P., Roman, J., Rosetti, M., Roy, A., Santos, R., Schauer, S., Schupback, R., Strauss, N., Subramanian, A., Talamas, J., Tesfaye, S., Theodore, J., Viel, R., Vo, A., Wilson, B., Wu, X., Wyman, D., Ye, W.J., Young, G., Zainoun, J., Zembek, L., Zimmer, A. and Zody, M. Direct Submission Submitted (11-DEC-2001) Whitehead Institute/MIT Center for Genome Research, 320 Charles Street, Cambridge, Ma 02141, USA On Dec 11, 2001 this sequence version replaced gi:15042917. All repeats were identified using RepeatMasker:	Choepel, Y., Colangelo, M., Collins, S., Collymore, A., Cook, A., Choepel, Y., Colangelo, M., Collins, S., Collymore, A., Cook, A., Cooke, P., DeArellano, K., Dewar, K., Diaz, J.S., Dodge, S., Faro, S., Ferreira, P., FitzHugh, W., Gage, D., Galagan, J., Gardyna, S., Ginde, S., Gord, S., Horton, L., Hulme, W., Iliev, I., Johnson, R., Jones, C., Kamat, A., Karatas, A., Kells, C., LaRocque, K., Lamazares, R., Landers, T., Lehoczky, J., Levine, R., Liu, G., MacLean, C., Macdonald, P., Major, J., Marquis, N., Matthews, C., Meneus, L., Mihova, T., Mlenga, V., Murphy, T., Naylor, J., Nguyen, C., Norbu, C., Norman, C.H., O'Connor, T., O'Donnell, P., O'Neil, D., Oliver, J., Peterson, K., Phunkhang, P., Pierre, N., Pollara, V., Rasman, J., Rosetti, M., Roy, A., Santos, R., Stange-Thomann, N., Seaman, S., Severy, P., Sougnez, C., Spencer, B., Stange-Thomann, N., Sesiltey, H., Viel, R., Vo, A., Wilson, B., W., X., Trigilio, J., Topham, K., Travers, M., Travis, N., Trigilio, J., Young, G., Zainoun, J., Zembek, L., Zimmer, A. and Zody, M. Direct Submission L. Submitted (31-JUL-2001) Whitehead Institute/MIT Center for Genome Research, 320 Charles Street, Cambridge, MA 02141, USA and erson, S., Barna, N., Bastien, V., Boguslavkiy, L., Boukhgalter, B., Anderson, S., Barna, N., Bastien, V., Boguslavkiy, L., Boukhgalter, B., Anderson, S., Barna, N., Bastien, V., Boguslavkiy, L., Boukhgalter, B.,	Straus, N., Subramanian, A., Talamas, J., Testaye, S., Theodore, J., Tristrell, A., Travers, M., Triglio, J., Vassiliev, H., Viel, R., Vo, A., Wilson, B., Wu, X., Wyman, D., Ye, W.J., Young, G., Zainoun, J., Zimmer, A. and Zody, M. Direct Submission Submitted (25-OCT-2000) Whitehead Institute/MIT Center for Genome Submitted (25-OCT-2000) Whitehead Institute/MIT Center for Genome Research, 320 Charles Street, Cambridge, MA 02141, USA 3 (bases 1 to 156235) Birren, B., Linton, L., Nusbaum, C., Lander, E., Ali, A., Allen, N., Barta, N., Basta, N., Bastien, V., Boguslavkiy, L., Boukhgalter, B., Roderson, S., Barna, N., Bastien, V., Boguslavkiy, L., Boukhgalter, B.,
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Query Match 90.0%; Score 18; DB 9; Length 156235; Best Local Similarity 100.0%; Pred. No. 2.1e+02; Matches 18; Conservative 0; Mismatches 0; Indels 0; Gaps

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       in cancer, particularly colon, breast and oral tumours, embryonic development disorders, angiogenic disorders and wound healing. Th antisense sequences, including the one shown here, can be used in
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                                 suppression of Focal Adhesion Kinase expression for the of cancers of the breast, colon, mouth or skin (especially a, and angiogenic disorders e.g. retinal neovascularisation -
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present invention relates to novel antisense compounds which can be for modulating the expression of human focal adhesion kinase (FAK)

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The antisense compounds comprise antisense oligonucleotides (8-30 nucleotides in length) targeted to the 5'-untranslated region, translational termination region or 3' untranslated region of a nucleic acid molecule encoding FAK. The antisense oligonucleotides can be used to inhibit the expression of FAK mRNA. The antisense oligonucleotides represent potential chemotherapeutic agents in the treatment of melanoma and the prevention of tumour metastasis. The antisense compounds and methods are useful for treating diseases associated with overexpression or constitutive activation of FAK. Such diseases include cancers of the breast, colon, mouth or skin (especially a melanoma), and anglogenic disorders such as retinal neovascularisation. AASI5383-AASI5422
The invention describes an isolated polynucleotide and polypeptide useful for stimulating and/or expanding T cells specific for a tumour protein for determining the presence of a cancer in a patient. A composition containing the polynucleotide and/or polypeptide is useful for treating a lung cancer in a patient. The polypeptide is useful for emoving tumour cells from a biological sample. The polynucleotide is
                                                                                                                             Novel polynucleotide encoding a lung stimulating and/or expanding T cells
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                                                                                                 Claim 1; SEQ ID No 981; 223pp; English.
                                                                                                                                                                                                                Marnerakis M,
                                                                                                                                                                                                                                                                                                   13-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                               10-JUL-2001; 2001WO-US22058
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200204514-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    represent the FAK antisense oligonucleotides of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ш
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         tumour; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TTTCAACCAGATGGTCATTC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TTTCAACCAGATGGTCATTC 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                          CORIXA CORP
                                                                                                                                                                                                Watanabe Y, Henderson RA, Jo
is M, Carter D, Fanger GR, V
.. Wang A, Fanger N, Switzer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard; cDNA; 550
                                                                                                                                                                                                                                                                                   2000US-0651563.

2000US-0658824.

2000US-0671325.

2000US-0677419.

2000US-0702705.

2000US-0703705.

2000US-07036457.

2001US-0849626.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                   2000US-0614124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    lung tumour protein clone R0127:G07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; 5 C; 3 G; 7 T; 0 other;
                                                                                                                           expanding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cell;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 20;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           response stimulator;
                                                                                                                           specific for a tumour protein
                                                                                                                                                                                                                Vedvick TS,
                                                                                                                                                                                                   ۶
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 2
                                                                                                                                                                                                 Mcneil1
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                                                                                                                                                                                                Retter MW;
Bangur CS
l PD, Clap
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                                                                                                                                                                                                Clapper
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8

(first entry)

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Antisense suppression of Focal Adhesion Kinase expression for the treatment of cancers of the breast, colon, mouth or skin (especially a melanoma), and angiogenic disorders e.g. retinal neovascularisation
                                                                                                                                                                                      Human; focal adhesion kinase, FAK; melanoma; tumour metastasis;
cancer; angiogenic disorder; retinal neovascularisation; cytostatic;
ophthalmological; ss.
                                                                                                                                                    Human focal adhesion kinase (FAK) cDNA seguence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 2; Page -; 19pp; English.
                      AAS15382/c
ID AAS15382 standard; cDNA; 3791 BP.
                                                                                                                                                                                                                                                                                                                                                                                                             19-AUG-1999; 99US-0377310.
13-JUL-2000; 2000WO-US18999.
                                                                                                                                                                                                                                                                                                                                                                           09-JAN-2001; 2001US-0757100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaarde WA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-010103/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MONI/) MONIA B P.
(GAAR/) GAARDE W A.
(NERO/) NERO P S.
                                                                                                                                                                                                                                                                                                   US2001034329-A1.
                                                                                                                                                                                                                                                                Homo sapiens,
                                                                                                                  29-JAN-2002
                                                                                                                                                                                                                                                                                                                                     25-OCT-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Monia BP,
                                                                              AAS15382;
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                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sequences to the human focal adhesion kinase (FAK) protein. This protein is involved in integrin-mediated signal transduction, and is implicated in cancer, particularly colon, breast and oral tumours, embryonic development disorders, angiogenic disorders and wound healing. The antisense sequences can be used in the treatment of all of these.
also useful as probe or primer to detect the level of mRNA encoding a tumour protein. This sequence encodes a lung tumour associated protein or protein fragment, described in the method of the invention.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human, focal adhesion kinase, FAK, signal transduction, cancer, embryonic development disorder, angiogenic disorder, wound healing; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes a number of phosphorothioate antisense
                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cancer
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New antisense compounds for inhibiting focal adhesion kinase expression, especially useful for inhibiting retinal neovascularization, or for diagnosing and treating e.g. colon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 20; DB 22; Length 3791;
Pred. No. 1.9;
                                                                                                                                                                                    100.0%; Score 20; DB 24; Length 550; 100.0%; Pred. No. 1.4;
                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 3791 BP; 1110 A; 849 C; 924 G; 908 T; 0 other;
                                                                                                                                                Sequence 550 BP; 155 A; 113 C; 121 G; 161 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human focal adhesion kinase coding sequence.
                                                                                                                                                                                                                          0; Mismatches
                                                                                                              ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Column 31-40; 30pp; English.
                                                                                                                                                                                                                                                                                                                                                                                        AAC65534 standard; DNA; 3791 BP
                                                                                                                                                                                                                                                                                195 TTTCAACCAGATGGTCATTC 214
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                                                                                                                                                                                                     Best Local Similarity 100.
Matches 20; Conservative
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Best Local Similarity 100.
Matches 20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaarde WA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-006141/01.
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                                                                                             rom WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US6133031-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-0CT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Monia BP,
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                                                                                                                                                                                      Query Match
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Nero PS;

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The present invention relates to novel antisense compounds which can be used for modulating the expression of human focal adhesion kinase (FAK). The antisense compounds comprise antisense oligonuclectides (8-30 nuclectides in length) targeted to the 5'-untranslated region, translational termination region or 3' untranslated region of a nucleic acid molecule encoding FAK. The antisense oligonuclectides can be used to inhibit the expression of FAK mRNA. The antisense oligonuclectides represent potential chemotherapeutic agents in the treatment of melanoma and the prevention of tumour metastasis. The antisense compounds and methods are useful for treating diseases associated with overexpression
                                                                                                                                                                                                                                                                                                                                                                                    breast, colon, mouth or skin (especially a melanoma), and angiogenic disorders such as retinal neovascularisation. The present cDNA sequence for human FAK is used to design the FAK antisense oligonucleotides
                                                                                                                                                                                                                                                                                                                                                       or constitutive activation of FAK. Such diseases include cancers of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Note: This sequence (Genbank accession number L13616) is not given in the patent but is acquired by the indexer from Genbank from the information provided in the patent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 3791 BP; 1110 A; 849 C; 924 G; 908 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; r. 0;
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Matches 20, Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  of the invention.
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ID ABV2919
XX
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RESULT 7
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Best Local (
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16-MAR-2000; 2000US-189862P.
25-MAY-2000; 2000US-207454P.
09-JUN-2000; 2000US-211314P.
18-JUL-2000; 2000US-219007P.
13-DEC-2000; 2000US-255281P.
                                                                                                                                                       3536
                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to an isolated nucleic acid molecule (I) compared a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the specification or its complement. (I) is useful for:

(a) assessing whether a patient is afflicted with prostate cancer;

(b) monitoring the progression of prostate cancer in a patient;

(c) assessing the efficacy of a test compound to inhibit prostate
             Human DNA sequence
                                     14-FEB-2002
                                                                                      AAS94837
                                                                                                                                                                                                                                                         Sequence 4052 BP; 1199 A; 906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, usef for detecting presence of prostate cancer, stage of prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-662795/76.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; prostate cancer; cytostati pharmacogenomic marker; gene; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABV29192;
                                                                                                                                                                                                                                                                                                                                                                                       cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-FEB-2001; 2001WO-US05171
                                                                                                                                                                                                                                                                                                           £6£6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MILL-)
                                                                                                                                                                                                                                                                                                                                                                         (d) assessing the efficacy of a therapy for inhibiting prostate cancer
                                                                                                                                                                                                                  Local Similarity
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                                                                                                                                                                                                                                                                                                                                                              b
                                                                                                                                                                                                                                                                                                         selecting a composition for inhibiting prostate cancer in a patient; assessing the prostate cell carcinogenic potential of a compound; determining whether prostate cancer has metastazed in a patient, assessing the aggressiveness or indolence of prostate cancer in a
                                                                                                                                                                   TTTCAACCAGATGGTCATTC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sapiens
                                                                                                                                                       TTTCAACCAGATGGTCATTC 3517
                                                                                                                                                                                                         20;
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                                                                                                                                                                                                                                                                                 also useful as a pharmacodyanamic or pharmacogenomic marker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MILLENNIUM PREDICTIVE MEDICINE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ,
R
                                                                                       standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Page 6216-6217; 11750pp; English.
                                                                                                                                                                                                                                                                                                                                                                                      patient,
                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first
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                                     (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             expression marker cDNA 29183
           #92 expressed during
                                                                                       DNA;
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                                      entry)
                                                                                                                                                                                                                  100.0%;
                                                                                       4481
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cytostatic; carcinogen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Monahan JE;
                                                                                                                                                                                                       0
                                                                                                                                                                                                                  Score 20; DB
Pred. No. 1.9;
                                                                                                                                                                                                                                                         C; 971 G;
                                                                                                                                                                                                          Mismatches
             foam cell differentiation
                                                                                                                                                                                                                                                          970 T; 6
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                                                                                                                                                                                                                              Length
                                                                                                                                                                                                                                                           other;
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                                                                                                                                                                                                                                   4052;
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RRESULT 8
AA1978A29
XX AA19
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DT 13-N
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequences that are differentially expressed during foam cell differentiation. The polynucleotide sequences of the invention or a composition comprising these polynucleotides are useful as a high throughput method for detecting altered expression of one or more polynucleotides in a sample. The polynucleotides can be used in the diagnosis of disorders associated with foam cell development such as atheroseclerosis, cerebral stroke, and cardiovascular disorders such a coronary artery disease. The polynucleotide sequences can also be used as PCR primers and probes. The polynucleotides of the invention are a useful in gene therapy. AAS94746-AAS95021 represent the human polynucleotide sequences of the invention which are differentially expressed during foam cell differentiation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3558
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            are
                                                                             02-MAR-2001;
                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                              Human neuroblastoma expressed polynucleotide SEQ ID NO 3903
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAI97828
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to the isolation of human sequences that are differentially expressed during foam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Composition useful for diagnosis of conditions, disorders or diseases associated with atherosclerosis, comprises several polynucleotides that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Shiffman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; foam cell differentiation; atherosclerosis; cerebral stroke; cardiovascular disorder; coronary artery disease; gene therapy; ds.
                                                                                                                                                                                      WO200166719-A1
                                                                                                                                                                                                                                                                                                                                                                                                     13-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-010925/01.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1; Page 145-146; 315pp; English.
                                                                                                                                                                                                                                                                                            neuroblastoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4481
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           standard; cDNA; 707
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            'n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                             2001WO-JP01629
                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Somogyi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 1326 A; 974 C; 1045 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%;
                                                                                                                                                                                                                                                                                         malignancy; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 20;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            in foam cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               æ
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                                                                                                                                                                                                                                                                                         marker; N-myc; TrkA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 4481;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          polynucleotide cell
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07-MAR-2000; 2000JP-0159195

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WO200157271-A2.
                                                                                                                                                                                                                                                                                                                      Homo sapiens.
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                                                                                                                                                                                                                                            01-FEB-2002
                                                                                                                                                                                                                                                                                                                                                             09-AUG-2001
                                                                                                                                                                                                                        ABA51579;
                                                                                   Query Match
                                                                                                          Matches
                                                                                                                                                                                RESULT 10
                                                                                                                                                                                         ABA51579
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                                                                                                                                             The invention relates to novel genes (AAI93926-AAI97963) expressed in human neuroblastoma. The nucleic acids are applicable as a probe or primer in diagnosing the prognosis of human neuroblastoma, malignancy and susceptibility indicators or tumour markers for anti-cancer agents. The gene information for diagnosing prognosis is related to factors similar to that for N-myc and TrkA genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         white rot basidiomycete, Coriolus versicolor. The present invention also describes a method for detecting at least one sulphur-containing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence encodes an aldo/ketoreductase (I) isolated from the white rot basidiomycete, Coriolus versicolor. The present invention also
                                                                     Nucleic acids originating in gene expressed in human neuroblastoma, useful as probe or primer in diagnosing prognosis of human neuroblastoma, malignancy and susceptibility indicator or tumour marker
                                                                                                                                                                                                                                                                                                                                                                                                                                            Aldo/ketoreductase; white-rot basidiomycete; methylbenzothiophene; AAD; aryl-alcohol dehydrogenase; enzyme; gene; ds.
                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                         Coriolus versicolor aldo/ketoreductase encoding DNA SEQ ID NO:1.
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                                                                                                                                                                                                                                          92.0%; Score 18.4; DB 22; Length 707; 95.0%; Pred. No. 9.8;
                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                      Sequence 707 BP; 200 A; 129 C; 149 G; 224 T; 5 other;
                                                                                                                                                                                                                                                                 ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /product= "aldo/ketoreductase"
                                                                                                                                                                                                                                                   Pred. No. 9.8; Mismatches
                                                                                                                          Claim 1; Page 2870-2871; 2979pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A gene encoding a new protein and its use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
354..1229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Page 7-8; 12pp; Japanese.
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        (HISM ) HISAMITSU PHARM CO LTD
                                                                                                                                                                                                                                                                                                        431 Trrcaaccaggregreatre 450
                                                                                                                                                                                                                                                                                    1 TTTCAACCAGATGGTCATTC 20
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                                                                                                                                                                                                                                                                                                                                                                                                     17-APR-2002 (first entry)
                                                                                                                                                                                                                                       Query Match
Best Local Similarity 95.0°
Matches 19; Conservative
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(CHIB-) CHIBA PREFECTURE
                                                                                                      for anti-cancer agents
                                                  WPI; 2001-565584/63.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Coriolus versicolor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-158675/21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (WARI/) WARIISHI H.
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                              Nakagawara A;
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                                                                                                                                                                                                                                                                                                                                                                                 ABA93211;
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The invention relates to a spatially-addressable set of single exon nucleic acid probes for measuring gene expression in a sample derived from human breast and BT 474 cells. The method involves conteacting the probes with a collection of detectably labelled nucleic acids derived from mRNA of human breast, and then measuring the label bound to each probe of the microarray. The probes are useful for verifying the expression of regions of genomic DNA predicted to encode proteins. They are useful for gene discovery, and for encode proteins. They are useful for gene discovery, and for determining predisposition and/or prognosing breast disease. Gene expression analysis is useful for assessing the toxicity of chemical agents on cells. The microarray of this invention presents a far greater diversity of probes for measuring gene expression, with far less bias than expressed sequence tag microarrays. The method is suitable for rapid production of functional information from genomic sequence. The present sequence is a single exon nucleic acid probe of the invention.
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hydrocarbon compound selected from methylbenzothiophenes in which, in a process of assaying a compound by the expression of (1), the increase in the expression of (1) shows that the compound is the objective compound (1) and the gene encoding (1) can be used for decomposing noxious
                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                    compounds
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; microarray; single exon probe; gene expression; breast;
                                                                                                                                                                                                                                                        DB 24; Length 1311;
58;
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                                                                                                                                                substances, particularly sulphur-containing heterocyclic
                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human breast cell single exon nucleic acid probe #10274.
                                                                                                                                                                                                                       Sequence 1311 BP; 340 A; 332 C; 391 G; 248 T; 0 other;
                                                                                                                                                                                                                                                                                                            100.0%; Preα. ...
've 0; Mismatches
                                                                                                                                                                                                                                                                                          Score 17;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABA51579 standard; DNA; 244
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21-SEP-2000; 2000US-0234687.
27-SEP-2000; 2000US-0236359.
04-OCT-2000; 2000GB-0024263.
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30-JUN-2000; 2000US-0608408
                                                                                                                                                                                                                                                                                          85.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            357 TTCAACCAGATGGTCAT 373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                   2 TTCAACCAGATGGTCAT 18
                                                                                                                                                                                                                                                                                                                                                                    17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    disease; cancer; ss.
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RESULT 11
ABA36555
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Query Match
Best Local Similarity
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                                                                                                   measuring human gene expression in a sample derived from human heart. present sequence is one such probe. The probes may be used for predicting, measuring and displaying gene expression in samples derive from the human heart via microarrays. By measuring gene expression, the probes are useful for predicting, diagnosing, grading, staging, monitoring and prognosing diseases of the human heart and vascular systems. Cardiovascular disease, hypertension, cardiac arrhythmias and
                                                                                                                                                                                                                                                                                                   Penn
                                  Sequence
                                                                   specification,
                                                                    Note: The sequence data for this patent did not form specification, but was obtained in electronic format
                                                                                          congenital heart disease.
                                                                                                                                                                                         The present invention relates to single exon nucleic acid probes for
                                                                                                                                                                                                                 Claim 4;
                                                                                                                                                                                                                                                   Single exon nucleic acid probes
                                                                                                                                                                                                                                                                            WPI; 2001-488899/53
                                                                                                                                                                                                                                                                                                                                                  04-OCT-2000;
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27-SEP-2000;
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26-MAY-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200157274-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         congenital heart
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; gene expression; heart; microarray; vascular system; probe; cardiovascular disease; hypertension; cardiac arrhythmia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Probe
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                                                                                                                                                                                                                                                                                                                          (MOLE-)
                                                                                                                                                                                                                                                                                                                                                                                                30-JUN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           118
                                                     ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 TTTCAACCAGATGGTCATTC 20
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                                                                                                                                                                                                                                                                                                                          MOLECULAR DYNAMICS INC
                                                                                                                                                                                                               SEQ ID No 15021; 530pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standard;
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                                                                                                                                                                                                                                                                                                                                               2000US-0180312
2000US-0207456
2000US-0608408
2000US-0632366
2000US-0234587
2000US-0236359
2000US-0236359
2000US-0236353
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                                  A; 59 C;
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90.0%;
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Score
Pred.
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Pred. No. 56;
                                  36 G;
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16.8;
No. 56;
                                  99 T; 0 other;
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            22;
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           Length
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                                                                    part of the printed directly from WIPO
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             244;
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RESULT 13
ABA46482
ID ABA46
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AC ABA46
XX
DT 01-FE
XX

ABA46482

standard;

DNA; 400

01-FEB-2002 ABA46482;

(first entry)

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RESULT 12
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                                          Matches
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                                                              Query Match
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                                                                                                        The present invention provides a number of single exon nucleic acid probes which are derived from genomic sequences expressed in the human bone marrow. They can be used to measure gene expression in bone marrow samples, which may enable the improved diagnosis and treatment of cancers such as lymphoma, leukaemia and myeloma. The present sequence is one of the probes of the invention.
                                                                                                                                                                                                                                                              Penn
                                                                                    Sequence 244
                                                                                                                                                                                   Example 4; SEQ ID NO:
                                                                                                                                                                                                                                                                                                                            03-AUG-2000;
21-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                            04-FEB-2000;
26-MAY-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; bone marrow expressed exon; gene expression analysis; probe; microarray; cancer; leukaemia; lymphoma; myeloma; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human bone marrow expressed single exon probe SEQ ID NO:
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                                                                                                                                                                                                                                                                                   (MOLE-)
118
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                                                                                                                                                                                                                                                              SG,
             TTTCAACCAGATGGTCATTC 20
TTTCAACCAGATGTACATTC 137
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                                                                                                                                                                                                         genome-derived single exon nucleic acid probes useful for 
zing gene expression in human bone marrow -
                                         l Similarity
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                                                                                                                                                                                                                                                                                   MOLECULAR DYNAMICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard; DNA;
                                                                                                                                                                                                                                                              Hanzel DK,
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                                                                                                                                                                                                                                                                                                                            2000US-0608408.
2000US-0632366.
2000US-0234687.
                                          Conservative
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2000US-0207456.
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                                                                                                                                                                                                                                                                                                         2000US-0236359.
2000GB-0024263.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                   50 A; 59 C;
                                                  84.0%;
90.0%;
                                                                                                                                                                                   18273; 658pp + Sequence Listing;
                                                                                                                                                                                                                                                              Chen W,
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                                                    Score 16.8;
Pred. No. 5
                                                                                    36 G;
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                                          Mismatches
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                                                                                    0 other;
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                                                               22;
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                                         Indels
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                                         Gaps
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measuring human gene expression in a sample derived from human heart. The present sequence is one such probe. The probes may be used for predicting, measuring and displaying gene expression in samples derived from the human heart via microarrays. By measuring gene expression, the probes are useful for predicting, diagnosing, grading, staging, monitoring and prognosing diseases of the human heart and vascular system e.g. cardiovascular disease, hypertension, cardiac arrhythmias and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Single exon nucleic acid probes for analyzing gene expression in human hearts -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to single exon nucleic acid probes for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; bone marrow expressed exon; gene expression analysis; probe; microarray; cancer; leukaemia; lymphoma; myeloma; ss.
     Human; gene expression; heart; microarray; vascular system; probe;
cardiovascular disease; hypertension; cardiac arrhythmia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human bone marrow expressed single exon probe SEQ ID NO: 5265
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    at ftp.wipo.int/pub/published_pct_sequences.
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0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; SEQ ID No 5154; 530pp; English.
                                                                                                                                                                                                                                                                                                                                                                                               Rank DR
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30-JUN-2000; 2000US-0608408.
03-AUG-2000; 2000US-0632366.
21-SEP-2000; 2000US-0234687.
27-SEP-2000; 2000US-0234587.
04-OCT-2000; 2000GB-0024263.
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                                        congenital heart disease; ss
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les 18; Conservative
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                                                                                                            WO200157274-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         from human breast and BT 474 cells. The method involves contacting the probes with a collection of detectably labelled muclaic acids derived from human breast, and then measuring the labelled muclaic acids derived from mRNA of human breast, and then measuring the label bound to each probe of the microarray. The probes are useful for verifying the expression of regions of genomic DNA predicted to determining predisposition and/or prognosing breast disease. Gene expression analysis is useful for assessing the toxicity of chemical expression analysis is useful for assessing the toxicity of chemical expression analysis is useful for assessing the toxicity of chemical expression analysis is useful for assessing the toxicity of chemical diversity of probes for measuring gene expression, with far less bias than expressed sequence tag microarrays. The method is suitable for rapid production of functional information from genomic sequence. The present sequence data for this patent did not form part of the invention. Note: The sequence data for this patent did not form part of the part of the printed specification, but was obtained in electronic format directly
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        invention relates to a spatially-addressable set of single exon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New spatially-addressable set of single exon nucleic acid probes, useful for measuring gene expression in sample derived from human breast, comprises number of single exon nucleic acid probes -
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                                    Human; microarray; single exon probe; gene expression; breast;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID NO 5177; 327pp + sequence listing; English.
Human breast cell single exon nucleic acid probe #5177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              from WIPO at ftp.wipo.int/pub/published_pct_sequences.
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2000US-0234687.
2000US-0236359.
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                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-496933/54
                                                        disease; cancer; ss
                                                                                                                         WO200157271-A2
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21-SEP-2000;
27-SEP-2000;
                                                                                         Homo sapiens
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26-MAY-2000;
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Search completed: January 12, 2003, 09:51:21 Job time : 255 secs
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Best Local Similarity 90.0
Matches 18; Conservative
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26-MAY-2000;
30-JUN-2000;
30-AUG-2000;
03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
                                                                                                                                                                                          The present invention provides a number of single exon nucleic acid probes which are derived from genomic sequences expressed in the human bone marrow. They can be used to measure gene expression in bone marrow samples, which may enable the improved diagnosis and treatment of cancers such as lymphoma, leukaemia and myeloma. The present sequence is one of the probes of the invention.
                                                                                                                                                              Sequence 400 BP; 94 A; 97 C; 84 G; 125 T; 0 other;
                                                                                                                                                                                                                                                                                                                   Penn
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                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-488900/53.
                                                                                                                                                                                                                                                                                                                                                                                                                   (MOLE-) MOLECULAR DYNAMICS INC.
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                                                         344 TTTCAACCAGATGTACATTC 363
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; 2000US-0207456.
; 2000US-0608408.
; 2000US-0632366.
; 2000US-0234687.
; 2000US-0236559.
; 2000US-0236559.
                                                                                                            84.0%; Score 16.8; DB 22; Length 400; 90.0%; Pred. No. 61; tive 0; Mismatches 2; Indels 0
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Appli Appl Appl Appli
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Sequence 1, Application US/09377310B
Sequence 1, Application US/09377310B
Setent No. 6133031
GENERAL INFORMATION:
APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310B
CURRENT APPLICATION NUMBER: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO!
LENGTH: 3791
                                                                                                        Sequence 2
Sequence 2
Sequence 2
Sequence 5
Sequence 9
Sequence 1
Sequence 1
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Patent NO. 6133031
GENERAL INFORMATION:
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GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENERAL GENE
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US-08-664-5968-4
US-08-721-798A-1
US-08-9461-527-75
US-08-724-394A-21
US-08-724-394A-21
US-08-724-394A-21
US-09-233-086-53
US-09-641-638-91
US-08-467-944A-1
US-08-467-947A-1
US-08-178-252-19
US-09-178-252-20
US-09-178-252-20
US-09-357-071-39
US-09-357-071-39
US-09-357-071-39
US-09-357-071-39
US-09-357-071-39
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: antisense sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Artificial Sequence
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/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
                                      GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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US-08-961-527-73
US-08-921-527-73
US-09-329-920-1
US-08-639-501-1
US-09-044-906-1
US-09-453-7028-57
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US-08-477-329-1
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US-08-980-400-1
US-09-583-459A-1
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US-09-583-459A-1
US-09-68-149-1
US-08-46-910-1
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Listing first 45 summaries
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seq length: 200000000
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Minimum DB Maximum DB

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                                                                                                                                                                                            APPLICATION NUMBER: US 07/146,877
PILING DATE: 22-JAA-1988
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/347,291
PILING DATE: 02-MAY-1989
ATTORNEY AGENT INFORMATION:
NAME: MAKI J.D., David J.
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 990008.4460
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PAGES: 823-830
DATE: 1993-11
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LOCATION: (233)..(3391)
PUBLICATION INFORMATION:
                                                                                             TELEFAX: 206-682-6031
TELEX: 3723836
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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VOLUME: 12
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Release #1.24 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
HYPOTHETICAL: N
                  MOLECULE TYPE:
                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: TITLE OF INVENTION:
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                                                                                                                                  TELEPHONE: 206-01
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                                             STRANDEDNESS:
                                                                TYPE:
                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: FILING DATE:
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                              TOPOLOGY:
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Kindsvogel Ph.D., Wayne R.

KINDSVOGEN PRODUCING SECRETED RECEPTOR ANALOGS
VENTION: METHODS OF PRODUCING SECRETED POLYPEPTIDE
VENTION: AND BIOLOGICALLY ACTIVE DIMERIZED POLYPEPTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sledziewski Ph.D., Andrzej Z
                                linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Seed and Berry
                                                                                                                                                                 206-622-4900
                  CDNA
                                                 double
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FUSIONS
                                                                                                                                                                                                                                                                                                                                                             US/07/634,510
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                                                                                                                                                                                               990008.446C3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 20; L
Pred. No. 0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           . 28;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 3791;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0,
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NAME/KEY: CDS;
LOCATION: 354..3671;
OTHER INFORMATION:
US-08-180-195-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 90.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 1, Application US/08477329 Patent No. 5750375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 206-682-6031
TELEX: 3723836
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                             MOLECULE I HYPOTHETICAL: I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.24
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1263 TITCATCCTGATGGTCATTC 1244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
ADDRESSEE: Seed and
                                                                                                                                                                    ANTI-SENSE: N
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION: TELEPHONE: 206-622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Maki, David J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ANTI-SENSE:
                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 4465 base pair
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Bell, Lillian A. APPLICANT: Kindsvogel Ph.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
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IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                               MOLECULE TYPE:
                             ORGANISM: Homo sapiens
DEVELOPMENTAL STAGE: Adult
TISSUE TYPE: Skin
CELL TYPE: fibroblasts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: ZUU CONTRACTOR CONTRAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/477,329
FILING DATE: 07-JUN-1995
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TISSUE TYPE: Skin
CELL TYPE: fibroblasts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 TTTCAACCAGATGGTCATTC 20
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                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                     1: 4465 base pairs nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Seattle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            E: Seed and Berry 6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                    double
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           METHODS OF PRODUCING SECRETED RECEPTOR ANALOGS AND BIOLOGICAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0,
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Pred. No. 1:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 4465;
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RESULT 6
US-08-980-400-1/c
US-08-980-400-1/c
Sequence 1, Application US/08980400
Fatent No. 6018026
Fatent No. 6018026
FAPPLICANT: Sledziewski Ph.D., Andrzej Z
APPLICANT: Bell, Lillian A.
FAPLICANT: Rindsvogel Ph.D., Wayne R.
TITLE OF INVENTION: METHODS OF PRODUCING SECRETED RECEPTOR ANALOGS AND BIOLOGICAL
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSEE: Seed and Berry
STREET: 6300 Columbia Center, 701 Fifth Avenue
CITY: Seattle
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  Gaps
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  Indels
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90.0%; Pred. No. 13;
iive 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIPCATION:
PRIOR APPLICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/477,329
FILING DATE: 07-UW-1995
ATTORNEY/AGENT INFORMATION:
NAME: MAKH, David J.
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 990008.446C6
TELEPHONE: 206-622-4900
TELEPHONE: 206-682-6031
  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA
ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.24
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/980,400
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                                                                              Db 1263 rrrcarccreardercarrc 1244
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DEVELOPMENTAL STAGE: Adult
TISSUE TYPE: Skin
                                          1 TTTCAACCAGATGGTCATTC 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 4465 base pairs
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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  Conservative
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Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: CDN.
HYPOTHETICAL: N
ANTI-SENSE: N
ORIGINAL SOURCE:
ORGANISM: Homo s.
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; LOCATION: 354..3
; OTHER INFORMATION
US-08-980-400-1
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IMMEDIATE SOURCE:
CLONE: pR-rX1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY:
18;
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Matches
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                                          ઠે
                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Sledziewski Ph.D., Andrzej Z
APPLICANT: Bell, Lillian A.
APPLICANT: Bell, Lillian A.
APPLICANT: Kindsvogel Ph.D., Wayne R.
TITLE OF INVENTION: METHODS OF PRODUCING SECRETED RECEPTOR ANALOGS AND BIOLOGICAL
CORRESPONDENCES: 36
CORRESPONDENCE ADDRESS:
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                                                                                                                                                              Gaps
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                                                                                                                      DB 1; Length 4465;
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                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AUDRESSEE: Seed and Berry STREET: 6300 Columbia Center, 701 Fifth Avenue CITY: Seattle
                                                                                                                                         ed. No. 13;
Mismatches
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Pred. No. 13;
                                                                                                                 Query Match

Best Local Similarity 90.0%; Pred. No. 13
Matches 18; Conservative 0; Mismatche
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ama: 990008.446D5
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ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
SOFTWARE: PATENTION DATA:
APPLICATION NUMBER: US/08/475,458
FILING DATE: 06-JUN-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: MAKI, David J.
REGISTRATION NUMBER: 990008.4461
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION ON 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 4465 base pairs
                                                                                                                                                                                                                                                                                                                                      Sequence 1, Application US/08475458 Patent No. 5843725 GENERAL INFORMATION:
                                                                                                                                                                                                                                          1263 TTTCATCCTGATGGTCATTC 1244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Homo sapiens
DEVELOPMENTAL STAGE: Adult
                                                                                                                                                                                                    1 TTTCAACCAGATGGTCATTC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          84.0%;
90.0%;
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CELL TYPE: fibroblasts
IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: CDNA
HYPOTHETICAL: N
ANTI-SENSE: N
              NAME/KEY: CDS
LOCATION: 354..36
COTHER INFORMATION:
US-08-477-329-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; LOCATION: 354..30
; OTHER INFORMATION
US-08-475-458-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORIGINAL SOURCE:
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US-08-475-458-1/c
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US-09-583-459A-1/c
                                                                                                              US-09-583-459A-1
                                   Query Match
Best Local Similarity 90.0
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 1, Application US/09583459A Patent No. 6291212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Sledziewski Ph.D., Andrzej Z
APPLICANT: Bell, Lillian A.
APPLICANT: Kindsvogel Ph.D., Wayne R.
TITLE OF INVENTION: METHODS OF PRODUCING
TITLE OF INVENTION: AND BIOLOGICALLY ACT
TITLE OF INVENTION: FUSIONS
                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 206-682-6031
TELEX: 3723836
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                        TOPOLUCE TYPE: C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 99
TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 02-MAY-1989
ATTORNEY/AGENT INFORMATION:
NAME: Maki J.D., David J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 27-DEC-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.24
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                               FEATURE:
                                                                                                                                                                                                                                                                                            ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                         HYPOTHETICAL:
ANTI-SENSE: N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: BOOK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                 MMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 4465 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: 200-682-6031
                                                                                                                           NAME/KEY: CDS
LOCATION: 354..3671
OTHER INFORMATION:
                                                                                                                                                                                                                            TISSUE TYPE: Skin
CELL TYPE: fibroblasts
                                                                                                                                                                                                                                           ORGANISM: Homo sapiens
DEVELOPMENTAL STAGE: Adult
TISSUE TYPE: Skin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 07/146,877 FILING DATE: 22-JAN-1988 APPLICATION NUMBER: US 07/347,291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: FILING DATE: 30-MA CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE:
1 TTTCAACCAGATGGTCATTC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98104-7092
                                                                                                                                                                                              pR-rX1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-MAY-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-DEC-1990
                                                                                                                                                                                                                                                                                                                                            CDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     David J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         METHODS OF PRODUCING SECRETED RECEPTOR ANALOGS AND BIOLOGICALLY ACTIVE DIMERIZED POLYPEPTIDE FUSIONS
                                                               84.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US 07/634,510
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                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        990008.446C3
                                                                 Score 16.8;
Pred. No. 13;
                                                Mismatches
                                                                              DB 4;
                                                                            Length 4465;
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                                                  Gaps
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US-09-583-210-1
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Patent No. 6291646
Query Match 84.0
Best Local Similarity 90.0
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 07/634,510
FILING DATE: 27-DEC-1990
APPLICATION NUMBER: US 07/146,877
FILING DATE: 22-JAN-1988
APPLICATION NUMBER: US 07/347,291
FILING DATE: 02-MAY-1989
ATTORNEY/AGENT INFORMATION:
NAME: Maki J.D., David J.
REGISTRATION NUMBER: 31,392
REGISTRATION NUMBER: 990008.446
                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 206-682-6031
TELEX: 3723836
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Sledziews...
APPLICANT: Bell, Lillian A.
APPLICANT: Bell, Lillian A.
APPLICANT: Kindsvogel Ph.D., Wayne R.
APPLICANT: Kindsvogel Ph.D., Wayne R.
TITLE OF INVENTION: METHODS OF PRODUCING SECRETED RECEPTOR ANALOGS
TITLE OF INVENTION: AND BIOLOGICALLY ACTIVE DIMERIZED POLYPEPTIDE
TITLE OF INVENTION: FUSIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1263 TTTCATCCTGATGGTCATTC 1244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.24
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/583,21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                         FEATURE:
                                                                                                                                                                                          IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                     ANTI-SENSE: N
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 4465 base pair
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION: TELEPHONE: 206-622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                          HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: cDNA
                                                                                            LOCATION: 354..36
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 3
CLASSIFICATION:
                                                                                                                NAME/KEY: CDS
LOCATION: 354..3671
                                                                                                                                                                                                              TISSUE TYPE: Skin
CELL TYPE: fibroblasts
                                                                                                                                                                         CLONE:
                                                                                                                                                                                                                                                 ORGANISM: Homo sapiens DEVELOPMENTAL STAGE: A
                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Seattle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98104-7092
                                                                                                                                                                                                                                                                                                                                                                                                     nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                    linear
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                                                                                                                                                                                                                                                                                                                                                                                     double
84.0%; Score 16.8; 1
90.0%; Pred. No. 13;
tive 0; Mismatches
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                                                                                                                                                                                                                                                     Adult
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         990008.446C3
                                       DB 4;
   2;
                                       Length 4465;
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   Gaps
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RESULT 10

US-09-433-059-1/c

| Sequence 1, Application US/09435059
| Sequence 1, Application US/09435059
| Patent No. 6323323
| GENERAL INFORMATION:
| APPLICANT: Sledation Sequence 1
| APPLICANT: Bell, Lillian A. APPLICANT: Kindsvogel Ph.D., Wayne R. TITLE OF INVENTION: METHODS OF PRODUCING SECRETED RECEPTOR ANALOGS AND BIOLOGICAL NUMBER OF SEQUENCES: 36
| CORRESPONDENCE ADDRESS: ADDRESSEE: Seed and Berry | STREET: 6300 Columbia Center, 701 Fifth Avenue | CITY: Seattle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURREATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.24
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/435,059
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/477,329
FILING DATE:
ATTORNEY/ACENT INFORMATION:
NAME: Maki, David J.
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 990008.446C6
TELEPHONE: 206-622-4900
TELEPAX: 206-682-6031
TELEPAX: 206-682-6031
TELEFX: 3723336
INFORMATION FOR SEG ID NO: 1:
SEQUENCE CHARRACTERISTICS:
LENGTH: 4465 base pairs
TYPE: nucleic acid
STRANDEDMESS: double
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                84.0%; Score 16.8; I
90.0%; Pred. No. 13;
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                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: WA
COUNTRY: USA
ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPACIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                              1263 TTTCATCCTGATGGTCATTC 1244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1263 TTTCATCCTGATGGTCATTC 1244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Homo sapiens
DEVELOPMENTAL STAGE: Adult
    1 TITCAACCAGAIGGICATIC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CELL TYPE: fibroblasts
IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 TTTCAACCAGATGGTCATTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 90.0°
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: CDNA
HYPOTHETICAL: N
ANTI-SENSE: N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; LOCATION: 354..30; OTHER INFORMATION US-09-435-059-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: CDS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  g
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                                                                                                                                                                                         APPLICANT: Slediewski Ph.D., Andrzej Z
APPLICANT: Slediewski Ph.D., Andrzej Z
APPLICANT: Bell, Lillian A.
APPLICANT: Bell, Lillian A.
TITLE CA.
TITLE OF INVENTION: METHODS OF PRODUCING SECRETED RECEPTOR ANALOGS
TITLE OF INVENTION: FUSIONS
TITLE OF INVENTION: FUSIONS
TITLE OF INVENTION: FUSIONS
AUTHER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSEE: Seed and Berry
STREET: Seatle
CITY: Seattle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: USA
ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYBE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.24
CURRENT APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 07/634,510
FILING DATE: 27-DEC-1990
APPLICATION NUMBER: US 07/146,877
FILING DATE: 22-JAN-1988
APPLICATION NUMBER: US 07/146,877
FILING DATE: 22-JAN-1989
ATTORNEY/AGENT INFORMATION:
NAME: MAKI J.D., DAVIG J.
NAME: MAKI J.D., DAVIG J.
NAME: MAKI J.D., DAVIG J.
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION OF SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 4465 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   990008.446C3
                                                                                                            US-09-583-449A-1/c
; Sequence 1, Application US/09583449A
Patent No. 6300099
; GENERAL INFORMATION:
1263 TTTCATCTGATGGTCATTC 1244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ANTI-SENSE: N
ORIGINAL SOURCE:
ORGANISM: Homo sapiens
DEVELOPMENTAL STAGE: Adult
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CELL TYPE: Skin
CELL TYPE: fibroblasts
IMMEDIATE SOURCE:
CLONE: PRATURP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: CDS
LOCATION: 354..3671
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: CDNA HYPOTHETICAL: N
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US-08-168-917-1/c
                                                                   US-08-460-510-1/c
                                                                                  RESULT 12
                                                                                                                                                                                                                                                              NAME/KEY:
;
LOCATION:
US-08-168-917-1
                Sequence 1, Application US/08460510 Patent No. 5872218 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1, App11
Patent No. 568657
                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEI.

NAME: Ching, Edwin F. 4,090
REGISTRATION NUMBER: 34,090
REFERENCE/DOCKET NUMBER: 1241f
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 136-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (415) 326-2422
INFORMATION FOR SEQ ID NO: 11
SEQUENCE CHARACTERISTICS:
LENGTH: 5427 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                   1096 TTTCATCCTGATGGTCATTC 1077
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                           ANTI-SENSE: NO ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Ching, Edwin P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Escobedo, Jaime A. APPLICANT: Williams, Lewis T.
                                                                                                                                                                                                                                                                                                                                                                               HYPOTHETICAL: F
                                                                                                                                                                                                                                                                                                                                                                                           TOPULCE TYPE: CU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                            ORGANISM: Homo Sapiens
STRAIN: lambda gt10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
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                                                                                                                                                                 TTTCAACCAGATGGTCATTC 20
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 Wolf, David
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Giese, Neill A.
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Tomlinson, James
                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                             linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                          double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Larry J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DOMAINS OF EXTRACELLULAR REGION OF HUMAN PLATELET-DERIVED GROWTH FACTOR RECEPTOR POLYPEPTIDES
                                                                                                                                                                                                               90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US/07/650,793
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                                                                                                                                                                                                               Score 16.8;
Pred. No. 1
                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                 DB 1;
                                                                                                                                                                                                                                Length 5427;
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RESULT 13
US-08-460-490-1/c
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                                                                                                                                                                                                       Sequence 1, Application US/08460490 Patent No. 5891652
                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                   GENERAL INFORMATION:

APPLICANT: Wolf, David

APPLICANT: Tomlinson, James E.

APPLICANT: Fretto, Larry J.

APPLICANT: Giese, Neill A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (415) 326-24: INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                             1096 TTTCATCCTGATGGTCATTC 1077
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Release #1.0, Version #1.
CURRENT APPLICATION DATA:
                                                 APPLICANT: Williams TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Williams
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 326-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INTI-SENSE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Dow, Karen B.
REGISTRATION NUMBER: 29,684
REFERENCE/DOCKET NUMBER: 01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                               Local Similarity es 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN: lambda gt10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET:
                                                                                                                                                                                                                                                                                                                                             1 TTTCAACCAGATGGTCATTC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        94105
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One Market Plaza, Steuart Street Tower, Suite 2000
                                                                                Escobedo, Jaime A. Williams, Lewis T.
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WILLIAMS, Lewis T.
WILLIAMS, LEWIS T.
WILLIAMS, LEWIS T.
WENTION: DOMAINS OF EXTRACELLULAR REGION OF HUMAN
JENTION: PLATELET-DERIVED GROWTH FACTOR RECEPTOR POLYPEPTIDES
                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Giese,
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Giese, Neill A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   E: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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187..3504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
TOWNSEND and TOWNSEND and CREW
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                                                                                                                                                                                                                                                                                                                                                                                               84.0%;
                                                   DOMAINS OF EXTRACELLULAR REGION OF HUMAN PLATELET-DERIVED GROWTH FACTOR RECEPTOR POLYPEPTIDES
                                   23
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Pred. No. 1:
                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                 DB 2;
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0; Mismatches
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                                FILING DATE: 10-FBB-1989

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/151,414
FILING DATE: 02-FBB-1988
ATTORNEY/AGENT INFORMATION:
NAME: Dow, Karen B.
REGISTRATION NUMBER: 29,684
REFERENCE/DOCKET NUMBER: 2307K-267-2
FLECOMMUNICATION INFORMATION:
TELEPHONE: 415/326-242
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 5427 base pairs
TYPE: nucleic cid
STRANDEDNESS: single
                 APPLICATION NUMBER: US 07/309,322
FILING DATE: 10-FEB-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 3, Application US/08461917; Patent No. 6372438
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IELECOMMUNICATION INFORMATION:
TELEPHONE: 415/326-2400
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Best Local Similarity 90.0
Matches 18; Conservative
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: CDS
LOCATION: 187..3507
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US-08-461-917-3/c
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Patent No. 6043211
GENERAL INFORMATION:
APPLICANT: ESCOBEDO, Jaime A.
TITLE OF INVENTION: PLATELET-DERIVED GROWTH FACTOR RECEPTORS
INVERSE OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew
STREET: One Market, Steuart Street Tower, 20th Floor
CITY: San Francisco
CITY: San Francisco
STATE: California
COUNTRY: USA
COUNTRY: USA
COUNTRY: USA
COUNTRY: WA
MEDIUM TYPE: Floppy disk
COMPUTER: Floppy disk
COMPUTER: Ploppy disk
COMPUTER: Ploppy disk
COMPUTER: PLOPS/MS-DOS
COUNTRY: STEEN PC COMPASIBLE FORM:
COMPUTER: PLOPS/MS-DOS
COMPUTER: PLOPS/MS-DOS
COMPUTER: PLOPS/MS-DOS
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: One Market Plaza, Steuart Street Tower, Suite 2000
San Francisco
California
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                                                                                                                                              MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: Patentin PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/460,490
FILING DATE: 02-UN-1995
CLASSITCATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: DOW, KARPE B.
REGISTRATION NUMBER: 012418-001420
TELEFROCH/DOCKET NUMBER: 012418-001420
TELEFROCHONICATION INFORMATION:
TELEFROCHONICATION INFORMATION:
TELEFROCH (415) 326-240
TELEFROCH (415) 326-242
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 5427 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
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APPLICATION NUMBER: US/08/462,728 FILING DATE: 05-JUN-1995 CLASSIFICATION: 435 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear
MOLECULE TYPE: CDNA to mRNA
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 TTTCAACCAGATGGTCATTC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Homo Sapiens
STRAIN: lambda gt10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 90.0
Matches 18; Conservative
                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: CDS
; LOCATION: 187..3504
US-08-460-490-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HYPOTHETICAL:
                                                                                                      94105
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Gaps
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APPLICANT: WILLIAMS, Lewis T.
APPLICANT: ESCOBEDO, Jaime A.
TITLE OF INVENTION:
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
DB 3; Length 5427;
                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Townsend and Townsend and Crew STREET: One Market, Steuart Street Tower, 20th Floor CITY: San Francisco STATE: California COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURBOTATION NUMBER: US/08/461,917
FILING DATE: 05-JUN-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: US 07/309,322
FILING DATE: 10-FEB-1989
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/309,322
FILING DATE: 02-FEB-1989
FILING DATE: 02-FEB-1988
ATFORDEY APPLICATION NUMBER: US 07/151,414
FILING DATE: 02-FEB-1988
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Sequence

Sequence 3191, App Sequence 242, App Sequence 180, App Sequence 181, App Sequence 171, App Sequence 171, App Sequence 28, Appl Sequence 212, Appl Sequence 112, Appl Sequence 112, Appl Sequence 2112, Appl

Sequence 3281 Sequence 1429

Seguence Sequence

Run on:

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Sequence 18, Application US/09757100B
; Sequence 18, Application US/09757100B
; Retent No. US20010034329A1
; GENERAL INFORMATION:
; APPLICANT: Hondia, Brett P.
; APPLICANT: Moria, Brett P.
; APPLICANT: Nero, Pamela S.
; TITLE OF INVENTION: Expression
; TITLE OF INVENTION: 1999-00-19
; PRIOR FILING DATE: 1999-00-19
; ROFTHARE: PETALING DATE: 2000-07-13
; NUMBER OF SEQ ID NOS: 44
; SOFTHARE: PETALING VEY: 2.0
; SEQ ID NO 18
                                                                                                                                                                                                                                                                                                                                US-09-925-300-712
US-09-954-456-2112
US-09-80-107-1609
US-09-864-761-21937
US-09-864-761-3281
US-09-864-761-3281
US-09-864-761-3281
US-09-864-761-3203
US-09-864-761-3203
US-09-815-242-9137

US-09-815-242-9537

US-09-938-842A-3191

US-09-938-842A-3191

US-09-974-300-088

US-09-974-300-180

US-09-974-300-180

US-09-974-300-180

US-09-974-300-180

US-09-974-300-208

US-09-974-300-208

US-09-101-368-171

US-09-170-62A-28

US-09-170-62A-28

US-09-170-62A-28

US-09-170-638-842A-975

US-09-170-8320-28

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100.0%; Pred. No. 0.13;
ive 0; Mismatches 0;
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Patent No. US20020168637A1
GPPERAL INFORMATION:
APPLICANT: Wang, Tongtong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
ORGANISM: Artificial Sequence
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      1 TTTCAACCAGATGGTCATTC
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US-09-736-457-981
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Best Local S:
Matches 20
    셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 15, Appl
Sequence 17, Appl
Sequence 19, Appl
Sequence 21, Appl
Sequence 1, Appli
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 18, Appl
Sequence 981, App
Sequence 981, App
                                                                                                                               January 12, 2003, 10:38:40; Search time 56 Seconds (without alignments) 157.180 Million cell updates/sec
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Sequence 1678
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1: \( \text{cgn2} \) \( \text{Fprodata} / 2 \) \( \text{pubpna} / \text{USO7} \) \( \text{PUBCOMB.seq:*} \) \( \text{cgn2} \) \( \text{Fprodata} / 2 \) \( \text{pubpna} / \text{PUST WW PUB.seq:*} \) \( \text{cgn2} \) \( \text{Fprodata} / 2 \) \( \text{pubpna} / \text{USO} \) \( \text{NEW PUB.seq:*} \) \( \text{cgn2} \) \( \text{Fprodata} / 2 \) \( \text{pubpna} / \text{USO} \) \( \text{NEW PUB.seq:*} \) \( \text{cgn2} \) \( \text{Fprodata} / 2 \) \( \text{pubpna} / \text{USO} \) \( \text{NEW PUB.seq:*} \) \( \text{cgn2} \) \( \text{Fprodata} / 2 \) \( \text{pubpna} / \text{USO8} \) \( \text{NEW PUB.seq:*} \) \( \text{cgn2} \) \( \text{Fprodata} / 2 \) \( \text{pubpna} / \text{USO9} \) \( \text{NEW PUB.seq:*} \) \( \text{cgn2} \) \( \text{Fprodata} / 2 \) \( \text{pubpna} / \text{USO9} \) \( \text{NEW PUB.seq:*} \) \( \text{cgn2} \) \( \text{Fprodata} / 2 \) \( \text{Pubpna} / \text{USO9} \) \( \text{NEW PUB.seq:*} \) \( \text{cgn2} \) \( \text{Fprodata} / 2 \) \( \text{Pubpna} / \text{USO9} \) \( \text{NEW PUB.seq:*} \) \( \text{12} \) \( \text{cgn2} \) \( \text{Fprodata} / 2 \) \( \text{Pubpna} / \text{USO} \) \( \text{NEW PUB.seq:*} \) \( \text{13} \) \( \text{cgn2} \) \( \text{Fprodata} / 2 \) \( \text{Pubpna} / \text{USO} \) \( \text{NEW PUB.seq:*} \) \( \text{13} \) \( \text{cgn2} \) \( \text{Fprodata} / 2 \) \( \text{Pubpna} / \text{USO} \) \( \text{NEW PUB.seq:*} \) \( \text{13} \) \( \text{cgn2} \) \( \text{Fprodata} / 2 \) \( \text{Pubpna} / \text{USO} \) \( \text{NEW PUB.seq:*} \) \( \text{13} \) \( \text{cgn2} \) \( \text{Fprodata} / 2 \) \( \text{Pubpna} / \text{USO} \) \( \text{PUBCOMB.seq:*} \) \( \text{13} \) \( \text{cgn2} \) \( \text{Fprodata} / 2 \) \( \text{Pubpna} / \text{USO} \) \( \text{PUBCOMB.seq:*} \) \( \text{13} \) \( \text{cgn2} \) \( \text{Fprodata} / 2 \) \( \text{Pubpna} / \text{USO} \) \( \text{PUBCOMB.seq:*} \) \( \text{14} \) \( \text{cgn2} \) \( \text{Fprodata} / 2 \) \( \text{Pubpna} / \text{USO} \) \( \text{PUBCOMB.seq:*} \) \( \text{14} \) \( \t
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                  GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                                   389086 seqs, 220051671 residues
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Maximum Match 100%
Listing first 45 summaries
                                                                                              OM nucleic - nucleic search, using sw model
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seq length: 200000000
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TITLE OF INVENTION: DIAGNOSIS OF LUNG CANCER FILE REFERENCE: 210121.478C15
CURRENT APPLICATION NUMBER: US/09/736,457
CURRENT FILING DATE: 2000-12-13
NUMBER OF SEQ ID NOS: 1864
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 981
LENGTH: 550
TYPE: DNA
ORGANISM: Homo sapien
US-09-736-457-981
US-09-849-626-981
; Sequence 981, Application US/09849626
; Publication No. US20020197669A1
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                                                            RESULT 4
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                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/902,941
CURRENT FILING DATE: 2001-07-10
NUMBER OF SEQ ID NOS: 2002
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 981
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 981, Application US/09902941 Patent No. US20020172952A1
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Best Local Similarity 100.0%;
Matches 20; Conservative 0
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APPLICANT: Vedvick, Thomas S.
APPLICANT: Bangur, Chaitanya S.
APPLICANT: Bangur, Chaitanya S.
APPLICANT: MCNabb, Andria
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210121.478C17
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                                                                                                                                                                                                                                                                  LENGTH: 550
TYPE: DNA
ORGANISM: Homo sapiens
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                                                                                                                                              1 TTTCAACCAGATGGTCATTC 20
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Watanabe, Yoshihiro
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Carter, Darrick
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Fan, Liqun
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                                                                                                                                                                                100.0%; Score 20; DB 9 ilarity 100.0%; Pred. No. 0.26; Conservative 0; Mismatches
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FILE REFERENCE: ISPH-0533
CURRENT APPLICATION NUMBER: US/09/757,100B
CURRENT FILING DATE: 2001-03-15
PRIOR APPLICATION NUMBER: 09/377,310
PRIOR FILING DATE: 1999-08-19
PRIOR APPLICATION NUMBER: PCT/US00/18999
PRIOR FILING DATE: 2000-07-13
NUMBER OF SEQ ID NOS: 44
SOFTMARE: PATENTIN VEY: 2.0
SEQ ID NO 1
SEQ ID NO 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; TYPE: DNA; ORGANISM: Homo sapien US-09-849-626-981
                                                                                                                                                                                                                                  NAME/KEY: CDS
LOCATION: (233)..(3391)
PUBLICATION INFORMATION:
JOURNAL: DNA
VOLUME: 12
ISSUE: 9
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                                                                       Matches
                                                                                        Query Match
Best Local Similarity
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LENGTH: 550
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APPLICANT: Gaarde, William A.
APPLICANT: Nero, Pamela S.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/849,626
CURRENT FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 1926
SOFTWARE: FastSEQ for Windows Version 3.0
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APPLICANT: McNeill, Patricia
APPLICANT: Clapper, Jonathan
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR
TITLE OF INVENTION: DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210121.478C16
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APPLICANT: Fanger, Gary
APPLICANT: Wang, Aijun
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                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                PÄGES: 823-830
DATE: 1993-11
DATABASE ACCESSION NUMBER: L13616/Genbank
DATABASE ENTRY DATE: 1995-01-02
                                                                                                                                                                                                                                                                                                                                                    FEATURE:
3529
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Wang, Aijun
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                                                                       Conservative
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Pred. No. 0.26;
                                                                       Mismatches
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US-864-761-5154

Sequence 5154, Application US/09864761

Sequence 5154, Application US/09864761

Sequence 5154, Application US/09864761

Sequence 5154, Application US/09864761

Septent No. US20020048763A1

SEPTENT SEPTENT HARDEN DAVIA REN

TITLE OF INVENTION: GENE REPRESSION ANALYSIS BY MICRORERAY

CURRENT APPLICATION NUMBER: 2001-05-28

FRIOR FILING DATE: 2001-05-28

FRIOR PELING DATE: 2000-05-26

FRIOR PELING DATE: 2000-05-27

FRIOR PELING DATE: 2001-05-27

FRIOR PELING DATE: 2001-05-27

FRIOR PELING DATE: 2001-05-27

FRIOR PELICATION NUMBER: COOL-10-10

FRIOR PELICATION NUMBER: COOL-10-10-10

FRIOR PELICATION NUMBER: C
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OTHER INFORMATION: EXPRESED IN BT474, SIGNAL = 0.97

OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 0.84

OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL
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SOFTWARE: Annomax Sequence I
SEQ ID NO 5154
LENGTH: 400
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ORGANISM: Homo sapiens
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Best Local Similarity
Matches 18; Conserv
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           Sequence 21875, Application US/09864761
Patent No. US20020048763A1
GENERAL INFORMATION:
APPLICANT: Penn, Sharron G.
APPLICANT: Rank, David R.
APPLICANT: Hanzel, David K.
APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
FILE REPERENCE: Application NUMBER: US/09/864,761
CURRENT APPLICATION NUMBER: US/09/864,761
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OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 0.84
OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 1
OTHER INFORMATION: SWISSPROT HIT: P47144, EVALUE 3.50e-02
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SEQ ID NO 21875
LENGTH: 244
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90.0%; Pred. No. 10;
iive 0; Mismatches 2;
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PRIOR PELLIANG DATE: 2010-02-04
PRIOR PELLICATION NUMBER: US 60/207,456
PRIOR PELLING DATE: 2000-05-26
PRIOR PELLING DATE: 2000-08-03
PRIOR PILING DATE: 2000-08-03
PRIOR PELLOCATION NUMBER: US 09/632,366
PRIOR PELLOCATION NUMBER: US 60/236,359
PRIOR PELLOCATION NUMBER: US 60/236,359
PRIOR PELLOCATION NUMBER: PCT/US01/00666
PRIOR PELLOCATION NUMBER: PCT/US01/00667
PRIOR PELLING DATE: 2001-01-30
PRIOR PELLOCATION NUMBER: PCT/US01/00667
PRIOR PELLOCATION NUMBER: PCT/US01/00669
PRIOR PELLING DATE: 2001-01-30
PRIOR PELLOCATION NUMBER: PCT/US01/00670
PRIOR PELLING DATE: 2001-01-30
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PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR FILING DATE: 2001-01-29
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OTHER INFORMATION: EXPRESSED IN BT47.
OTHER INFORMATION: EXPRESSED IN HEAR
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Matches 18; Conservative
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; OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL =
; OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL
US-09-864-761-16204
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PRIOR APPLICATION NUMBER: US 60/207,456
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 2000-05-26
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                                                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                     ENGTH: 528
 344
                                                                                            Local Similarity
                                     1 TTTCAACCAGATGGTCATTC 20
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FILING DATE: 2000-09-27
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FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00669
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TTTCAACCAGATGTACATTC 363
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                                                                            Conservative
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Pred. No. 12
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RESULT 9

US-09-866-510-15/c ; Sequence 15, Application US/09866510 ; Patent No. US20020111304A1

RESULT 11

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; LOCATION: (1)..(1683)
US-09-866-510-23
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Best Local Similarity
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                                                                    Matches
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                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/866,510
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/250,747
PRIOR FILING DATE: 2000-12-01
PRIOR PPLICATION NUMBER: 60/289,103
PRIOR FILING DATE: 2001-05-07
NUMBER OF SEQ ID NOS: 33
                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: KAZLAUSKAS, ANDRIUS
APPLICANT: IKUNO, YASUSHI
TITLE OF INVENTION: METHODS AND CÔMPOSITIONS FOR TREATING FIBROTIC DISEASES
FILE REFERENCE: ERM-104.01
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NUMBER OF SEQ ID NOS: 33
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CURRENT APPLICATION NUMBER: US/09/866,510
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/250,747
PRIOR FILING DATE: 2000-12-01
PRIOR APPLICATION NUMBER: 60/289,103
                                                                                                                                                                                                                                                                             SOFTWARE:
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TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR TREATING FIBROTIC DISEASES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: KAZLAUSKAS, ANDRIUS
APPLICANT: IKUNO, YASUSHI
                                                                                                                                                     NAME/KEY: CDS
LOCATION: (1)..(3270)
                                                                                                                                                                                        ORGANISM: Homo sapiens FEATURE:
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 910 TTTCATCCTGATGGTCATTC 891
                                                                                    Local
                                  1 TTTCAACCAGATGGTCATTC 20
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                                                                  l Similarity
18; Conserv
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                                                                  Conservative
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                                                                  Score 16.8; I
Pred. No. 18;
0; Mismatches
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Pred. No. 1
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Sequence 21, Application US/09866510;
Patent No. US2002011304A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: KAZLAUSKAS, ANDRIUS
APPLICANT: KAZLAUSKAS, ANDRIUS
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR TREATING FIBROTIC DISEASES
FILE REFERENCE: ERM-104-105-25
CURRENT APPLICATION NUMBER: 00/2-05-25
PRIOR FILING DATE: 2000-12-01
PRIOR FILING DATE: 2000-12-01
PRIOR FILING DATE: 2001-05-07
NUMBER OF SEQ ID NOS: 33
SOUTHANDER: PATENTION NUMBER: 60/289,103
PRIOR FILING DATE: 2001-05-07
SOUTHANDER: PATENTING UNCE: 2.1
INVENTION: METHODS AND COMPOSITIONS FOR TREATING FIBROTIC DISEASES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-955-953-1/C
; Sequence 1, Application US/09955363
; Patent No. US20020173621A1
; GENERAL INFORMATION:
; APPLICANT: Sledziewski Ph.D., Andrzej Z
; Kindsvogel Ph.D., Wayne R.
; Kindsvogel Ph.D., Wayne R.
; TITLE OF INVENTION: METHODS OF PRODUCING SECRETED RECEPTOR ANALOGS
; TITLE OF INVENTION: METHODS OF PRODUCING SECRETED POLYPEPTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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90.0%; Pred. No. 18;
iive 0; Mismatches
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Best Local Similarity 90.0%; Pred. No. 18;
Matches 18; Conservative 0; Mismatches
                  FILE REFERENCE: ERM-104.01
CURRENT APPLICATION NUMBER: US/09/866,510
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/250,747
PRIOR FILING DATE: 2000-12-01
PRIOR FILING DATE: 2000-12-01
PRIOR FILING DATE: 2001-05-07
NUMBER: OF SEQ ID NOS: 33
SEQ ID NO 19
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Matches 18; Conservative
                                                                                                                                                                                                                                                                       TYPE: DNA ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                         ) NAME/KEY: CDS
; LOCATION: (1)..(3318)
US-09-866-510-19
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US-09-866-510-21/c
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, LOCATION: (1)
US-09-866-510-21
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US-09-955-363-1/c
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                                   APPLICANT: IKUNO, YASUSHI
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR TREATING FIBROTIC DISEASES
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR TREATING FIBROTIC DISEASES
CURRENT APPLICATION NUMBER: US/09/866,510
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/280,747
PRIOR APPLICATION NUMBER: 60/289,103
PRIOR PILING DATE: 2000-12-01
PRIOR FILING DATE: 2000-05-07
NUMBER OF SEC ID NOS: 33
SOFTWARE: Patentin Ver. 2.1
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APPLICANT: KAZLAUSKAS, ANDRIUS
APPLICANT: KAZLAUSKAS, ANDRIUS
TITLE OF INVENTION METHODS AND COMPOSITIONS FOR TREATING FIBROTIC DISEASES
FILE REFERENCE: ERM-104.01
CURRENT APPLICATION NUMBER: 06/255
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 66/250,747
PRIOR PILING DATE: 2000-12-01
PRIOR APPLICATION NUMBER: 66/289,103
PRIOR PILING DATE: 2001-05-07
NUMBER OF SEQ ID NOS: 31
SOFTWARE: PATENTIN Ver. 2.1
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Best Local Similarity 90.0%; Pred. No. 18
Matches 18; Conservative 0; Mismatche
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US-09-866-510-17/c
; Sequence 17, Application US/09866510
; Patent No. US20020111304A1
; GENERALION:
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GENERAL INFORMATION:
APPLICANT: KAZLAUSKAS, ANDRIUS
APPLICANT: IKUNO, YASUSHI
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ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                    ; NAME/KEY: CDS
; LOCATION: (1)..(3318)
US-09-866-510-15
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; LOCATION: (1)..(3318)
US-09-866-510-17
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LENGTH: 3321
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STATE: WA
COUNTRY: USA
ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.24
CURRENT APPLICATION NUMBER: US/09/955,363
FILING DATE: 18-Sep-2001
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 07/634,510
FILING DATE: CUNKNOWN>
APPLICATION NUMBER: US 07/347,291
FILING DATE: CO-MAY-1989
ATTORNEY/AGENT INFORMATION:
NAME: MAKI J.D., DAVId J.
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 31,392
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 990008.446C3
TELEFAX: 206-622-4900
TELEFAX: 206-622-4900
TELEFAX: 206-622-4900
TELEFAX: 206-622-4900
TELEFAX: 3733836
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 4465 Dase pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: CDNA
HYPOTHETICAL: N
ANTI-SENSE: N_
                                                                                                                                                                                                                                          ; INAME/KEY: CDS
; LOCATION: 354..3671
; OTHER INFORMATION:
; SEQUENCE DESCRIPTION: SEQ ID NO: 1:
US-09-955-363-1
Search completed: January 12, 2003, 12:08:34 Job time : 59 secs
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                                                                                 1263 TITCATCCTGATGGTCATTC 1244
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ORIGINAL SOURCE:
ORGANISM: Homo sapiens
DEVELOPMENTAL STAGE: Adult
TISSUE TYPE: Skin
CELL TYPE: fibroblasts
IMMEDIATE SOURCE:
                                                                                                                         1 TTTCAACCAGATGGTCATTC 20
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Seed and Berry
STREET: 6300 Columbia Center
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STATE: WA
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AA65479 2x47a00.00
BK60918 600120623
AG060293 RC2-BN003
AG060293 Pan trog1
BF61197 602247296
AA922847 RC2-BN003
AQ424326 CITBI-E1-
BF771452 602149140
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BF217100 601813031
BF65659 602579335
BF217100 601813031
BF676992 602084209
BF217100 601813031
BF689361 601310527
BES66855 601339461
AN136503 UI-H-BII-
AA983514 ort1g10.s
BN664187 UI-E-CII-
N14230 yw7lh06.r1
A1141506 gx32212.x
BF084847 PM2-DT004
BG986106 PM2-DT004
A1016118 oct34907.s
AA404694 Zt41a11.r
A1380178 tf98e02.x
BM790849 K-EST0070
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CMO-GN0100-081100-576-a05 GN0100 Homo sapiens cDNA, mRNA sequence.
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20202663
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Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010,
Braail
Fil +55-11-2704922
Fax: +55-11-2707001
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                                          AI341506
BF084847
BG986106
AI016118
AA404694
AA1380178
BM790849
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BF669338
AW992843
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AQ424326
BF701452
BE701452
BE567556
BQ438629
AV726076
BE380127
BF565639
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BE617048
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BG110386
BG617438
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BF691197
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BF217100
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BF997449.1 GI:12403772
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 RESULT 1
BF997449/c
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DEFINITION
ACCESSION
VERSION
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AUTHORS
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MEDLINE
COMMENT
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SOURCE
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BE092864 CM2-BT074
R89170 yq02b06.r1
AV724065 AV724065
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(without alignments)
162.524 Million cell updates/sec
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         GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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                                                                                                                                                         16154066 seqs, 8097743376 residues
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                                           OM nucleic - nucleic search, using sw model
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gb_est5:*
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R32551/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20;
                                                                                                                                                     Email: est@watson.wustl.edu
Insert Size: 691
High quality sequence stops: 299
Source: IMAGE Consortium, LLNL
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Insert Length: 691
Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                    .M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,J. Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P. and Wilson,R. The WashU-Merck EST Project Unpublished (1995)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?t1=CMO&t2=CMO-GNO100-
081100-576-a05&t3=2000-11-08&t4=1)
                                                                                                                                       Seq
                                                                                                                                                                                                                                                                                                                                                                    Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R32551 351 bp mRNA linear EST 28-APR-199
yh54d07.r1 Soares placenta Nb2HP Homo sapiens cDNA clone
MAGE:133549 5' similar to gb:L05186 FOCAL ADHESION KINASE (HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             High quality sequence start: 10 High quality sequence stop: 285.
                                                                                       High quality sequence stop: 299
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                         Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hillier, L., Clark, N., Dubuque, T., Elliston, K.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/db_xref="GDB:539275"
/db_xref="taxon:9606"
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/dev_stage="Adult"
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                                             organism="Homo sapiens"
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Pred. No. 21;
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Best Local
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TTTCAACCAGATGGTCATTC 154
                            TTTCAACCAGATGGTCATTC 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20;
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                                                                                                                                                                                                                                                                                                                                            Contact: Zeguang Han
Chinese National Human Genome Center at Shanghai
351 Guo Shoujing Road, Zhangjiang Hi-Tech Park, Pudong, Shanghai
201203, P. R. China
Tel: 86-21-50801919 (ex.45)
Fax: 86-21-50801922
                                                                                                                                                                                                                                                                                                                                                                                                                                     Peng,Y., Song,H., Huang,Q., Huang,C., Gu,Y., Yang,Y., Gao,H., Xu,X., Li,N., Qian,B., Liu,F., Qu,J., Gao,X., Cheng, Zeng,L., Xu,S., Gu,W., Tu,Y., Jia,J., Fu,G., Ren,S., Zhon,G., Hu,R., Chen,J., Chen,Z. and Han,Z. Homo sapiens cDNA ADC clones
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 392)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AV709841 ADC Homo sapiens cDNA clone ADCAGAll 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                     Email: hanzg@chgc.sh.cn
This clone is available at C
Location/Qualifiers
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                                                                                                                                                                                 /tissue_type="Adrenal gland"
/dev_stage="Adult"
/lab_host="SOLR"
                                                                                                                                                                                                                                           db_xref="taxon:9606"
clone="ADCAGA11"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone_lib="Soares placenta Nb2HP"
/sex="Female"
                                                                                                                                                                                                                                                                          organism="Homo sapiens"
                                                                                                                                                                   note="Vector: pBluescript sk(-); Site_1: EcoRI; Site_2:
                                                                                                                                                                                                                              clone_lib="ADC"
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Pred. No. 24;
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                                                                                        Length 392;
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ACCESSION VERSION KEYWORDS SOURCE ORGANISM

AUTHORS

REFERENCE

MEDLINE COMMENT

JOURNAL

TITLE

DEFINITION

BE092864 LOCUS

RESULT 4

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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi; Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi; Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.

I (bases 1 to 416)

S Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M., Holman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,J., Rifkin,L., Rohlfing,T., Sorses,M., Tan,F., Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P. and Wilson,R.

AL Walliamson,A., Wohldmann,P. and Wilson,R.

Contact: Wilson RK

Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108

Tel: 314 286 1800

Email: est@watson.wustl.edu
Insert Size: 1206

High quality sequence stops: 337

Source: IMAGE Consortium, LLNL
This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.

Insert Length: 1206 Std Error: 0.00

Seq primer: M13RP1

High quality sequence stop: 337.

Location/Qualifiers

Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 424)
Gu, Y., Peng, Y., Song, H., Huang, Q., Yang, Y., Gao, G., Xiao, H., Xu, X., Li, N., Qian, B., Liu, F., Qu, J., Gao, X., Cheng, Z., Xu, Z., Zeng, L., Xu, S., Gu, W., Tu, Y., Jia, J., Fu, G., Ren, S., Zhong, M., Lu, G., Hu, R., Chen, Z., and Han, Z.
Homo sapiens cDNA HTE clones
Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Pac I and cloned into the Pac and Eco RI sites of the modified pT7T3 vector. Library went through one round of normalization. Library constructed by Bento Soares and M.Fatima Bonaldo."
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AV724065
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Homo sapiens"
/db_xref="GDB:3764781"
/db_xref="taxon:9606"
/clone="IMAGE:195731"
/clone="IMAGE:195731"
/sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 416;
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(A) Aradia = Homo sapiens"

(A) Arafe="taxon.9606"

(Clone_lib="BT0741"

(A) Stage="Adult"

(A) Stage="Adult"

(A) Note="Organ: Press; Vector: puc18; Site_1: Smal; Site_2: Smal; A mint-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196

716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and CDNA amplification were performed under low
                                                                                                                                                                                                                                                               Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (Bases I to 412)

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Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F.,

Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H.,

Brunstein,A., deoliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare

Simpson,A.J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tel: +55-11-2704922

Fax: +55-11-2707001

Email: asingbon@ludwig.org.br

This sequence was farived from the FAPESP/LICR Human Cancer Genome Project. This entry can be seen in the following URL (http://www.ludwig.org.br/scripts/gethtml2.pl?tl=&t2=CM2-BT0741-050 400-140-c128xs1=2000-04-05&t4=1)

Seq primer: puc 18 forward High quality sequence start: 39
High quality sequence stop: 410.
                                                                                    BE092864 12-JUN-2000 CM2-BT0741-050400-140-c12 BT0741 Homo sapiens cDNA, mRNA sequence.
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Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Shotgun sequencing of the human transcriptome with ORF expressed
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Proc. Natl. Acad. Sci. U.S.A. 97 (7), 3491-3496 (2000)
20202663
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Pred. No. 26;
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R89170
R89170.1 GI:953997
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Chinese National Human Genome Center at Shanghai
351 Guo Shoujing Road, Zhangjiang Hi-Tech Park, Pudong, Shanghai
201203, P. R. China
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
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BE566855.1 GI:9810575
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601339461F1 NIH_MGC_53 Homo
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Fax: 86-21-50801922
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Plate: LLCM364 row: i column: 15
High quality sequence stop: 421.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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This clone is available at CHGC in Shanghai.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CDNA Library Preparation: CLONTECH Laboratories, Inc. CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.gov
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/organism="Homo sapiens"
/db xref="taxon:9606"
/clone="IMAGE:3681878"
/clone="IMAGE:3681878"
/clone="IMAGE:3681878"
/clone="IMAGE:3681878"
/clone="IMAGE:3681878"
/clone="DH10B (T1 phage-resistant)"
/lab_host="DH10B (T1 phage-resistant)"
/note="Organ: bladder; Vector: pDNR-LIB (Clontech);
/note-"Organ: bladder; Vector: pDNR-LIB (
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/tissue_type="Hypothalamus"
/dev_stage="Adult"
/lab_host="SOLR"
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Pred. No. 26;
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IMAGE:3681878 5',
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ORIGIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TTTCAACCAGATGGTCATTC 357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TTTCAACCAGATGGTCATTC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The sequence contained an oligo-dT track that was present in the oligonuclectide that was used to prime the synthesis of first strand cDNA and therefore this may represent a bonafide poly A tail. cDNA Library Preparation: M.B. Soares Lab Clone distribution: NCI-CCAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: www-bio.lln1.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AW136503 429 bp mRNA linear EST 29-OCT-UI-H-BII-acx-e-10-0-UI.sl NCI_CGAP_Sub3 Homo sapiens cDNA clone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tumor Gene Index
Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa;
Mammalia; Eutheria;
1 (bases 1 to 429)
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AW136503.1 GI:6140563
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Seq primer: M13 Forward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
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Similarity 100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        117 a
/clone_lib="IMAGE:2715763"
/clone_lib="IMAGE:2715763"
/clone_lib="NCI_CGAP_Sub3"
/lab_host="Ph10B (Life Technologies)"
/lab_host="Ph10B (Life Tech
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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118 c 91 g 99 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               organism="Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chordata; Craniata; Vertebrata; Euteleostomi; Primates; Catarrhini; Hominidae; Homo.
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Pred. No. :
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Gaps

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BASE COUNT ORIGIN

DEFINITION

ACCESSION VERSION KEYWORDS

RESULT 9 AA983514

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ORGANISM

SOURCE

AUTHORS TITLE

JOURNAL

COMMENT

FEATURES

REFERENCE

BASE COUNT ORIGIN

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/closue_type="RPE and Choroid"
/dev_stage="adult" and Choroid"
/dev_stage="adult" and Choroid"
/lab_host="blublos (Life Technologies) (Tl phage resistant)"
/lab_host="blublos (Life Technologies) (Tl phage resistant)"
/note="Organ: eye; Vector: pT773-pec (Pharmacia) with a modified polylinker; Site_1: EcoR I; Site_2: Not I;
UI-E-CII is a normalized CDNA library containing the following tissue(s): RPE and Choroid. The library was constructed according to Bonaldo, Lennon and Soares,
Genome Research, 6:791-806, 1996. First strand CDNA synthesis was primed with an oligo-dT primer containing a Not I site. Double stranded cDNA was ligated to an EcoR I adaptor, digested with Not I; and cloned directionally into pT7T3-Pac vector. The oligonucleotide used to prime the synthesis of first-strand cDNA contains a library tag sequence that is located between the Not I site and the (dT)18 tail. The sequence etag for this library is ACCTA.
This library was created for the program, Gene Discovery in the Visual System, supported by National Eye Institute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Trugian for fact the Discovery and Mapping University of Iowa 451 Eckstein Medical Research Building Iowa City, IA 52242, USA Tel: 319 315 9265
Fax: 319 315 9266
Fax: 319 315 9265
Email: msoares@blue.weeg.uiowa.edu
Tissue Procurement: Dr. Gregg Hageman
CDNA Library preparation: Dr. M. Bento Soares, Univeristy of Iowa CDNA Library Arrayed by: Dr. M. Bento Soares, Univeristy of Iowa DNA Sequencing by: Dr. M. Bento Soares, Univeristy of Iowa Clone Distribution: Researchers may obtain clones from Research Genetics (www.resgen.com).
The following repetitive elements were found in this cDNA sequence: 1-22, AT_rich#Low_complexity (matched compliment)
Seg primer: 1-21, AT_rich#Low_complexity (matched compliment)
                                                                                                                                                                                                                                                                                                                                                           BM664187 linear EST 27-FEB-2002
UI-E-CI1-age-d-10-0-UI.s1 UI-E-CI1 Homo sapiens cDNA clone
UI-E-CI1-age-d-10-0-UI 3', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 (bases 1 to 447)
Bonaldo,M.F., Lennon,G. and Soares,M.B.
Normalization and subtraction: two approaches to facilitate gene
             100.0%; Score 20; DB 9; Length 432; 100.0%; Pred. No. 27; or Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Soares, MB
Program for Rat Gene Discovery and Mapping
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Homo sapiens"
/db xref="taxon:9606"
/clone="UI-E-CII-age-d-10-0-UI"
/clone=lib="UI-E-CII"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TAG_LIB=UI-E-CI1
TAG_TISSUE=RPE and Choroid
TAG_SEQ=ACCTA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Genome Res. 6 (9), 791-806 (1996)
97044477
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                                                                                                                                                                            91 TTTCAACCAGATGGTCATTC 110
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                                                                                                                                              1 TITCAACCAGAIGGICATIC
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Matches 20; Conserv
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ORGANISM
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AUTHORS
TITLE
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I Unpublished (1997)

Contact: Robert Strausberg, Ph.D.

Email: cgapbs-remail.nih.gov

Tissue Procurement: Christopher A. Moskaluk, M.D., Ph.D., Michael

Email: cgapbs-remail.nih.gov

Tissue Procurement: Christopher A. Moskaluk, M.D., Ph.D., Michael

Emmert-Buck, M.D., Ph.D.

CDNA Library Preparation: M. Bento Soares, Ph.D.

CDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be

found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llhl.gov/bbrp//amage/image.html

Insert Length: 1419 Std Error: 0.00

Seq primer: -40m13 fwd. ET from Amersham.
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/db_xref="taxon:9606"
/clone="IMAGE:1598466"
/clone="IMAGE:1598466"
/clone="taxon:960ed germ cell tumors"
/tissue_type="pooled_germ cell tumors"
/lab.host="Uhl10B"
/note="Vector: pT713D-Pac (Pharmacia) with a modified
polylinker; 1st strand cDNA was prepared from 3 pooled
germ cell tumors, and was then primed with a Not I -
oligo(dT) primer. Double-stranded cDNA was ligated to Eco
RI adaptors (Pharmacia), digested with Not I and cloned
into the Not I and Eco RI sites of the modified pT7T3
vector. Library is not normalized. Library was
constructed by Bento Soares and M. Fatima Bonaldo. "
3716-3720, 3733-3735 (INAGE CloneIDs 1257096-1258631, 1469064-1470883, 147552-1476743); NCI CGAP Pr22 pool 1 LLAM 2457-2459, 2758-2759, 3062-306 [TAMGE CloneIDs 985608-986759, 1101192-1101959, 1217928-1220615); NCI CGAP_CO10 pool 1 LLAM 2644-2653, 2871-2872 (INAGE CloneIDs 1057416-1061255, 1144584-1145351). Subtraction was performed as previously described [Bonnaldo, Lennon & Soares (1996): Normalization and Subtraction: Two Approaches To Facilitate Gene Discovery. Genome Research 6, 791-806.
TAG_IIB=NCI CGAP_GC4
TAG_TISSUE=Gene Cell
TAG_CAPARTC"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EST 27-AUG-1998
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NCI-CAAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 20; DB 10; Length 429; 100.0%; Pred. No. 27; cive 0; Mismatches 0; Indels 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                 131
                                                                                                                                                                                                                                                                                                                                                                                                                          105 g
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Best Local Similarity 100.0
Matches 20; Conservative
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TITLE
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Best Local Similarity
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               193
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                                                                                               Local
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             TTTCAACCAGATGGTCATTC
                                            TTTCAACCAGATGGTCATTC 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  High quality sequence stops: 330 Source: IMAGE Consortium, LLNL free through LLNL; contact the This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. 1 Tel: 314 286 1800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Tre,R., Williamson,A., Wohldmann,P. and Wilson,R. The WashU-Merck EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N42230 462 bp mRNA linear EST 24-JAN-19
yw71h06.rl Soares placenta 8to9weeks 2NbHP8to9W Homo sapiens cDNI
clone IMAGE:257723 5' similar to gb:L05186 FOCAL ADHESION KINASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tel: 314 286 1800 Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (1995)
Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hillier, L., Clark, N.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                               Similarity
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                                                                                                                                                                137
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Location/Qualifiers
                                                                             Conservative
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                                                                                                                                                             double-stranded cDNA was size selected, ligated to Eco RI adapters (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of a modified pT7T3 vector (Pharmacia). Library constructed by Bento Soares and M.Fatima Bonaldo."

1 96 c 89 g 139 t 1 others
                                                                                                                                                                                                                                                               /clone_lib="Soares_placenta_8to9weeks_2NbHP8to9W"
/dev_stage="two placentae: one from 8 weeks and another
from 9 weeks post conception"
/lab_host="DH10B (ampicillin resistant)"
                                                                                                                                                                                                                                                                                                                                                                                                  /db_xref="GDB:3887333"
/db_xref="taxon:9606"
/clone="IMAGE:257723"
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tive 0;
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ing,T., Soares,M., Tan,F., Trevaskis,E., Waterston
                 174
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                                                                             Score 20; DB 1
Pred. No. 28;
0; Mismatches
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Pred. No. 27;
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TTTCAACCAGATGGTCATTC 111
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Homo sapiens
Eukaryota; Metazoa; (
Mammalia; Eutheria; I
1 (bases 1 to 492)
                                                                                                                                                                                                                                                                                                                                                                                                                     20;
                                                                                                                                                             PM2-DT0042-100900-003-g03 DT0042 Homo sapiens BF084847
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome
Clone distribution: NCI-CGAP clone distribution
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 686 Std Error: 0.00
Seq primer: -40UP from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher A. Moskaluk, M.D.,
R. Emmert-Buck, M.D., Ph.D.
cDNA Library Preparation: M. Bento Soares, Ph.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         qx92a12.x1
similar to
                                                                                                                                       BF084847.1
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Unpublished (1997)
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note="Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site 1: Not I; Site 2: Ecc RI; Plasmid DNA from the normalized library NCI CGAP GC4 was prepared, and so circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-mplified CDNAs from a pool of 5,000 clones made from the same library (cloneIDs 1257096-1258631, 1469064-1470983, and 1475592-1476743). Subtraction by Bento Soares and M. Fatima Bonaldo. " a 98 c 123 g 132 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone="IMAGE:2009950"
(clone lib="NCI CGAP GC6"
/tissue_type="pooled germ cell tumors"
lab_host="DH10B"
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NCI CGAP GC6 Homo sapiens cDNA clone IMAGE:2009950 3'
gb:L05186 FOCAL ADHESION KINASE (HUMAN);, mRNA sequence.
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                       Chordata; Craniata; Vertebrata; Euteleostomi; Primates; Catarrhini; Hominidae; Homo.
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Pred. No. 28;
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KEYWORDS
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                                                                                                                                                                                                                                                                                                  Fax: +55-11-2707001

Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
This sequence was derived from the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=&t2=PM2-DT0042-100
900-003-g03&t5=2000-09-10&t4=1)
Seq primer: puc 18 forward
High quality sequence start: 5
High quality sequence stop: 492.
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РМ2-DT0042-190201-005-g03 DT0042 Homo sapiens cDNA, mRNA sequence.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 492)
Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R., Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F.,
Brunscain,A., deOliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
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Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,
Brazil
                                                                                                                         Shotgun sequencing of the human transcriptome with ORF expressed
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Laboratory of Cancer Genetics
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/organism="Homo sapiens"
/db xref="texon:9606"
/clone_ib="D40042"
/dev stage="Addut"
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Site 2: Sma1; A min1-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research)
profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and CDNA amplification were performed under low stringency conditions."
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Email: cgapbs-r@mail.nih.gov
cDNA Library Preparation: M. Bento Soares, Ph.D., M. Fatima Bonaldo
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DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 752 Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham
High quality sequence stop: 344.
                                                                                                                                                                        Fax: +55-11-2707001
Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=PM2&t2=PM2-DT0042-
190201-005-903&t3=2001-02-19&t4=1)
Seq primer: puc 18 forward
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1 (bases 1 to 527)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP), Tumor Gene Index
Unpublished (1997)
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,
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